Access DB#	
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### SEARCH REQUEST FORM

### Scientific and Technical Information Center

Requester's Full Na	ime:	Examiner # :  Serial Number:  Results Format Professor de	Date:
Mail Boy and Didas	Phone Number 30	Serial Number:	
	Room Location:	Serial Number: Results Format Preferred (circ	ele): PAPER DISK E-M
If more than one s	earch is submitted, please i	orioritize searches in order of	2.00 E W
*****	*********	***********************	need.
Include the elected spec utility of the invention.	les or structures kommende	ns. acronyms, and registry numbers, ar	subject matter to be searched
Title of Invention: _			
Inventors (please prov	ide full names):		
	1		· .
Earliest Priority Filin	ng Date:		
*For Sequence Searches	Only* Please include all pertinent info	rmation (parent, child, divisional, or issue	
appropriate serial number.	,	rmation (parent, child, divisional, or issue	d patent numbers) along with th
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	Bibliographic	Dr.Link	
c Completed:		Lexis/Nexis	
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rical Prep Time:	. Patent Family	WWW/Internet	
ine Time:	92 Other	Other (specify)	

PTO-1590 (8-01)



## STIC Search Report Biotech-Chem Library

#### STIC Database Tracking Number: 119583

TO: Terra Gibbs

Location: rem/2d10/2c18

Art Unit: 1635

Friday, April 16, 2004

Case Serial Number: 09/954556

From: David Schreiber

**Location: Biotech-Chem Library** 

Remsen E01A61 Phone: 272-2526

david.schreiber@uspto.gov

Search Notes	
	**





# STIC SEARCH RESULTS FEEDBACK FORM

#### Biotech-Chem Library

Questions about the scope or the results of the search? Contact the searcher or contact:

Mary Hale, Information Branch Supervisor Remsen Bldg. 01 D86 571-272-2507

Vol	untary Results Feedback Folin
>	I am an examiner in Workgroup: Example: 1610
>	Relevant prior art found, search results used as follows:
	☐ 102 rejection
	☐ 103 rejection
	☐ Cited as being of interest.
	Helped examiner better understand the invention.
	☐ Helped examiner better understand the state of the art in their technology.
	Types of relevant prior art found:
	☐ Foreign Patent(s)
	<ul> <li>Non-Patent Literature</li> <li>(journal articles, conference proceedings, new product announcements etc.)</li> </ul>
۶	Relevant prior art not found:
	Results verified the lack of relevant prior art (helped determine patentability).
	Results were not useful in determining patentability or understanding the invention.
Cor	nments:

Drop off or send completed forms to STIC-Biotech-Chem Library Remsen Bldg.



119583

#### Schreiber, David

From:

Gibbs, Terra

Sent:

Wednesday, March 31, 2004 4:20 PM

To:

Schreiber, David

Subject:

Sequence searcj request...

Hi David,

I have another request for a score over length search:

I need a length limited nucleotide sequence search against nucleobases 1479 through 1508 of SEQ ID NO:3 in USSN 09,954,556, where the returns are rank ordered based on the score over length/ratio as we've discussed. I need the lengths limited to hits between 8 and 50 nucleotides, and I'll take as many hits as you can import into excel (64,000?), and alignments for anything above .75 on the above ratio. Hope this is clear, please call me if it's not. I also need the interference databases searched if possible.

Terra Cotta Gibbs, Ph.D. Art Unit 1635 Remsen Building 2D10 571-272-0758

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1 (bases 1 to 17)

Karn, J., Gait M.J., Heaphy, S. and Dingwall, C. Viral (HIV) growth inhibition

Patent: US 6114109-A 17 05-SEP-2000;

Location/Qualifiers
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Karn, J., Gair, M. John., Heaphy, S. and Dingwall, C. RNA oligonucleotides that bind HIV tat protein Patent: US 5821046-A 20 13-OCT-1998;
Location/Qualifiers
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1 (Dases 1 to 18)

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Karn, J., Gait, M.John., Heaphy, S. and Dingwall, C. RNA oligonucleotides that bind HIV tat protein Patent: US 5821046-A 23 13-OCT-1998;
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/mcl_type="unassigned DNA"
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Unclassified.
Unclassified.
1 (bases 1 to 18).
1 (bases 1 to 18).
Karn, J., Gait, M.J., Heaphy, S. and Dingwall, C.
Viral (HIV) growth inhibition
Patent: US 6141109-A 23 05-SEP-2000;
Patent: US 6141109-A 23 05-SEP-2000;
Patent: Uscation/Qualifiers
18
                                                                  VIRAL (HIV) GROWTH INHIBITION
Patent: WO 9202228-A 17 20-FEB-1992;
Location/Qualifiers
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synthetic construct
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/mol_type="unassigned RNA"
/db_xref="taxon:32630"
                     /organism="synthetic construct"
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PR 05-JUN-1998 GB 9812196.5,02-MAR-1999 GB 9904790.4 PI
JONATHAN KARN, CATHERINE DENISE PRESCOTT
PC C12Q1/68,C12N15/09,G01N21/78,G01N33/53,G01N33/542,G01N33/566,
PC G1N37/00,
PC C12N15/00
CC Methods and Literary
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Sequence 1
AR340497
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Karn, J. and Prescott, C.D.

Methods and kits for discovery of RNA-binding Patent: US 6573045-A 1 03-JUN-2003;

Location/Qualifiers
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Human immunodeficiency virus 1
Viruses; Retroid viruses; Retroviridae; Lentivirus; Primate
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Methods and kits for discovery of RNA-binding Patent: JP 2002526032-A 1 20-AUG-2002;
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JP 2002526032-A/
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/mol_type="genomic RNA"
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                                      'mol_type="unassigned RNA"
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Human immunodeficiency virus
Viruses; Retroid viruses; Retroviridae; Lentivirus; Primate
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Solanum tuberosum
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AX325661
AX325661.1 GI:18096420
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Location/Qualifiers
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Patent: WO 0192512-A 1799 06-DEC-2001;
                                                                                                                                                                                                                                      Targeted chromosomal genomic alterations in
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/mol_type="unassigned DNA"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   /organism="Human immunodeficiency virus"
/mol_type="unassigned RNA"
/db_xref="taxon:12721"
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88.2%;
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Pred. No. 11;
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Best Local Similarity Matches 15; Conserv
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AX325662
AX325662.1
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Targeted Ciromosomal genomic alterations
single stranded oligonucleotides
Patent: WO 0192512-A 1800 06-DEC-2001;
UNIVERSITY OF DELAWARE (US)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta; Spermatophyta; Magnoliophyta; eudicotyledons; core eudicots; asterids; lamiids; Solanales; Solanaceae; Solanum.
1 (bases 1 to 18)
Karn, J., Gait, M. John., Heaphy, S. and Dingwall, C.
RNA oligonucleotides that bind HIV tat protein
                                                                                                                              Sequence 13
AR048072
                                                                                                                                                                                                                                                                                                                                                                                                                              VIRAL (HIV) GROWTH INHIBITION
Patent: WO 9202228-A 20 20-FEB-1992;
Location/Qualifiers
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synthetic construct
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                                                    Unclassified.
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1 (bases 1 to 18)
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1800 from Patent WO0192512.
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/mcl_type="unassigned RNA"
/db_xref="taxon:32630"
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/mol_type="unassigned DNA"
/db_xref="taxon:4113"
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Best Local Similarity 88.3
Matches 15; Conservative
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           RS Karn, J. and Prescott, C.D.

Methods and kits for discovery of RNA-binding compounds

AL Patent: JP 2002526032-A 3 20-AUG-2002;

RIBOTARGETS LTD

OS HIV

PN JP 2002526032-A/3

PD 20-AUG-2002

PF 04-UN-1999 GB 9812196.5,02-MAR-1999 GB 9904790.4 PI

JONATHAN KARN, CATHERINE DENISE PRESCOTT

PC C1201/68, C12N15/09, G01N21/78, G01N33/53, G01N33/542, G01N33/566,

PC G01N37/00,

PC C12N15/00

CC Methods and kits for discovery of RNA-binding compounds FH

Key

Location/Qualifiers

FT source 1. 18

FT source / Organism='HIV'.
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Unclassified.
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Location/Qualifiers
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/organism='HIV'
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/mol_type="unassigned DNA"
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/mol_type="unassign
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AR340499
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Methods and kits for discovery of rna-binding
Patent: WO 9964628-A 3 16-DEC-1999;
RIBOTARGETS LIMITED (GB)
                                                                                                                                                                                                                 Human immunodeficiency virus
Human immunodeficiency virus
Viruses; Retroid viruses; Re
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15; Conserv
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Karn, J. and Prescott, C.D.

Methods and kits for discovery of RNA-binding compounds
Patent: US 6573045-A 3 03-JUN-2003;
Location/Qualifiers
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                    AGCCAGACTTCAGCAGC 1506
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/mol_type="genomic RNA"
/db_xref="taxon:11676"
                                                                                      /organism="Human immunodeficiency
/mol_type="unassigned RNA"
/db_xref="taxon:12721"
                                                                                                                                                                                                                                                                                                                                                                                                                                               /mol_type="unassigned
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Sequence 18 from patent US
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Nucleic acids and secreted polypeptides encoded thereby
Patent: WO 0110902-A 40 15-FEB-2001;
Curagen Corporation (US)
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Unclassified.

1 (bases 1 to
                     Unknown.
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Location/Qualifiers
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/mol_type="unassigned DNA"
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Karn, J., Galt, M. John., Heaphy, S. and Dingwall, C. RNA oligonucleotides that bind HIV tat protein Patent: US 5821046-A 19 13-OCT-1998;
Location/Qualifiers
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RNA oligonucleotides that bind HIV tat protein
Patent: US 5821046-A 18 13-OCT-1998;
Location/Qualifiers
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Karn, J., Gair, M.John., Heaphy, S. and Dingwall, C. RNA oligonucleotides that bind HIV tat protein Patent: US 5821046-A 21 13-OCT-1998;

Location/Qualifiers
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/moi_type="unassigned DNA"
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Unclassified.
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1 (bases 1 to 17)
Karn,J., Gait,M.J., Heaphy,S. and Dingwall,C.
Viral (HIV) growth inhibition
Patent: US 6114109-A 18 05-SEP-2000;
Location/Qualifiers
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/mol_type="unassigned DNA"
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BD209410/c
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AR108983
Enzymatic nucleic acid treatment to hepatitis C virus infection. BD209410
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Patent: US 6114109-A 22 05-SEP-2000;
Location/Qualifiers
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Karn, J., Gait, M.J., Heaphy, S. and Dingwall, C. Viral (HIV) growth inhibition

Patent: US 6114109-A 21 05-SEP-2000;

Location/Qualifiers
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1 (bases 1 to 14)

AUTHORS

Blatt,L., McSwiggen,J.A., Roberts,E., Pavco,P.A. and Mack Enzymatic nucleic acid treatment of diseases or condition to hepatitis C virus infection Patent: JP 2002512791-A 3000 08-MAY-2002;

RIBOZYME PHARMACEUTICALS INC

OS Hepatitis virus (hepatitis C virus)

PD 08-MAY-2002

PP 26-APR-1999 JP 2000545991

PR 27-APR-1999 US 60/083217,18-SEP-1998 US 60/100842

25-FEB-1999 US 60/257608,23-MAR-1999 US 09/274553 PI

LAWRENCE BLATT,JAMES A MCSWIGGEN,ELISABETH ROBERTS,PAMELA A P. PAVCO,

PI DENNIS MACEJAK

PC C12N9/00,A61K31/7105,A61K38/21,A61K40''

PC A61K37/66,

PC C12N15/00

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Location/Qualifiers
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                                                                                                     /organism="synthetic construct"
/mol_type="unassigned DNA"
/db_xref="taxon:32630"
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1 (bases 1 to 14)

Brysch, W. and Schlingensiepen, K.
AN ANTISENSE OLIGONUCLECTIDE PREPARATION METHOD
Patent: WO 9833904-A 1715 06-AUG-1998;
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Brysch, W. and Schlingensiepen, K.

ANTISENSE OLIGONUCLECTIDE PREPARATION METHOD PAtent: WO 9833904-A 1714 06-AUG-1998;

BIOGNOSTIK GES (DE); BRYSCH WOLFGANG (DE)
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BD067079
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BD067080
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Schlingensiepen, K.H. and Brysch, W.
TITLE
An antisense oligonucleotide preparation method
Patent: JP 2001511000-A 1714 07-AUG-2001;
BICCNOSTIK GESELLSCHAFT FUR BICMOLEKULARE DIAGNOSTIK MBH
OS Unknown
OS Unknown
PN JP 2001511000-A/1714
PD 07-AUG-2001
PP 30-JAW-1998 JP 1998532533
PR 31-JAW-1997 EP 97101577
PI KARL HERMANN SCHTT
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An antisense oligonucleotide p
BD067080
BD067080.1 GI:22612683
JP 2001511000-A/1715.
unidentified
unidentified
1 (bases 1 to 14)
Schlingensiepen, K.H. and Brysch, W.
An antisense oligonucleotide preparation method
Patent: JP 2001511000-A 1715 07-AUG-2001;
BIOGNOSTIK GESELLSCHAFT FUR BIOMOLEKULARE DIAGNOSTIK MBH
OS Unknown
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An antisense oligonucleotide
BD067079.1 GI:22612682
JP 2001511000-A/1714.
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An antisense oligonucleotide preparation method Location/Qualifiers
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/mol_type="genomic_DNA"
/db_xref="taxon:32644"
                                                                                                                                                                                                                                                                                                                                                /organism='Unknown'
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/db_xref="taxon:32644"
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100.0%; Pred. No. 26;
Live 0; Mismatches
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                                  Query Match
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Matches 12; Conserv
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PC A
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JP 2002512791-A/2984
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Key Location/On-
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                                                                              /organism="unidentified"
/mol_type="genomic RNA"
/db_xref="taxon:32644"
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/mol_type="genomic DNA"
/db_xref="taxon:32644"
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                                  Score 10.8; D
Pred. No. 28;
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1 (bases 1 to 14)

Blatt, L., Mcswiggen, J.A., Roberts, E., Pavco, P.A. and Macejak, D.

Enzymatic nucleic acid treatment of diseases or conditions related
to hepatitis C virus infection

Patent: JP 2002512791-A 2984 08-MAY-2002;

RIBOZYME PHARMACEUTICALS INC
OS Hepatitis virus (hepatitis C virus)
PN JP 2002512791-A/2984
PD 08-MAY-2002
                                                                                                                                                                                                                                                                                                                                                                                                 PF 26-APR-1999 JP 2000545991
PR 27-APR-1998 US 60/083217,18-SEP-1998 US 60/100842 PR 25-PEB-1999 US 09/257608,23-MAR-1999 US 09/274553 PI LAWRENCE BLATT, JAMES A MCSWIGGEN, ELISABETH ROBERTS, PAMELA A PI
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07-AUG-2001
30-JAN-1998 JP 1998532533
31-JAN-1997 EP 97101531.8
KARL HERMANN SCHLINGENSIEPEN, WOLFGANG BRYSCH
C12N15/11, C07H21/04, A61K31/70
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19-JUN-1998 US 60/09097,19-JUN-1998 US 60/090079 PR
19-JUN-1998 US 60/090935,19-JUN-1998 US 60/089993 PR
19-JUN-1998 US 60/089992,19-JUN-1998 US 60/090072 PR
19-JUN-1998 US 60/089978,19-JUN-1998 US 60/090072 PR
19-JUN-1998 US 60/089991,19-JUN-1998 US 60/090043 PR
19-JUN-1998 US 60/090042,19-JUN-1998 US 60/090043 PR
19-JUN-1998 US 60/090044,19-JUN-1998 US 60/090043 PR
19-JUN-1998 US 60/090044,19-JUN-1998 US 60/090036 PR
19-JUN-1998 US 60/090047,19-JUN-1998 US 60/090037 PR
19-JUN-1998 US 60/090078,19-JUN-1998 US 60/090047 PR
19-JUN-1998 US 60/090078,19-JUN-1998 US 60/090047 PR
19-JUN-1998 US 60/090076,19-JUN-1998 US 60/090047 PR
19-JUN-1998 US 60/090076,19-JUN-1998 US 60/090047 PR
19-JUN-1998 US 60/090076,19-JUN-1998 US 60/090047 PR
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19-JUN-1998 US 60/090079,19-JUN-1998 US 60/090047 PR
19-JUN-1998 US 60/090079 PR
19-JUN-1998 US 60/0900 PR
19-JUN-1998 US 60/0900 PR
19-JUN-1998 US 60/0900 
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unidentified unidentified unclassified.
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1 (bases 1 to 10)

Roberts, B. L. and Shankara, S.

Preparation and use of superior vaccines

Patent: JP 2002534056-A 1327 15-OCT-2002;
                                                                                Sequence 72 from Pat
A71513
A71513.1 GI:4775125
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Preparation and use of superior vaccines
Key
Location/Qualifiers
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JP 2002534056-A/1327
15-OCT-2002
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72 from Patent WO9813521.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            /organism='Homo sapiens (human)'.
                                                                                                                                                                                                                                                                                                                                                                                                              /organism="Homo sapiens"
/mol_type="genomic DNA"
/db_xref="taxon:9606"
                                                                                                                                                                                                                                                                                                                                                      100.0%;
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AX623670
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AX003113.1
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Fesce, R. and Consalez, G.
METHOD FOR THE DIFFERENTIAL SCREENING OF GENE EXPRESSION
PRIMED REVERSE TRANSCRIPTION-POLYMERASE CHAIN REACTION
PALENT: WO 9813521-A 72 02-APR-1998;
FESCE RICCARDO (IT)
                                                                     Method for determining homeostasis of the skin Patent: WO 02053774-A 711 11-JUL-2002; Henkel Kommanditgesellschaft auf Aktien (DE) Location/Qualifiers
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Location/Qualifiers
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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/mol_type="unassigned
/db_xref="taxon:9606"
                                orcanism="Homo sapiens"
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AX629909
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Local Similarity 90.9%;
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalla; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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AX629388
AX629388.1 GI:28457426
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Location/Qualifiers
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae;
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                                                                                                   /organism="Homo sapiens"
/mol_type="unassigned DNA"
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/db_xref="taxon:9606"
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90.9%;
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RESULT 44

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TITLE
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Matches 10; Conserv
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Unclassified.
1 (bases 1 to 1
Inoue, T.
                                                                               Sequence 65
AR167701
AR167701.1
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AX631091
AX631091.1 GI:28459135
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Sequence 7143 from Patent
AX630102
                                          Unknown
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                                                        Unknown.
                                                                                                                        AR167701
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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                                                                                                                                                                                                                                                                                               /organism="Homo sapiens"
/mol_type="unassigned DNA"
/db_xref="taxon:9606"
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/mol_type="unassigned DNA"
/db_xref="taxon:9606"
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                                                                                  GI:17903498
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E29585/c
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E38691/c
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Best Local Similarity
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                                                                                                                                                                                   11 CCAAGAGGCCA 1
                                                                                                                                                                                                                                                                                                                                                                                                                    Method for amplifying DNA fragment, method for estimating state of microorganism existing and method for estimating state of microorganism existing and method for estimating state of waste Patent: JP 199276176-A 65 12-OCT-1999;
SANYO ELECTRIC CO LITD, SOCIETY FOR TECHNO-INNOVATION OF AGRICULTURE FORESTRY AND FISHERIES
OS Unidentified
PN JP 199276176-A/65
PD 12-OCT-1999
PF 31-WAR-1998 JP 1998087652
PR GOICHI INOUE
PC C12015/09,B09B3/00,C12Q1/00,C12Q1/68,C12N15/00,B09B3/00 CC
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12 bp DNA linear PAT 18-JUN-2001
Method for amplifying DNA fragment, method for estimating state of
microorganism existing and method for estimating state of waste.
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                                      unidentified
                                                    E38691.1 GI:18621353
JP 2000270867-A/65.
                                                                               E38691 12 bp DNA lir
Method and device for amplifying DNA fragment
E38691
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           unclassified.
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                                                                                                                                                                                                                                                                                             /organism="unidentified"
/mol_type="genomic DNA"
/db_xref="taxon:32644"
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Pred. No. 41;
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Query Match
Best Local Similarity
                                                                                                                                                                             Method for amplifying DNA fragment, amplification apparatus of DNA fragment, method for assaying a group of microorganisms, method for analyzing a group of microorganisms, method for analyzing a group of microorganisms, and method for assaying contaminating substance

AL Patent: JP 199341989-A 65 14-DEC-1999;
SANYO ELECTRIC CO LTT), SOCIETY FOR TECHNO-INNOVATION OF AGRICULTURE FORESTRY AND FISHERIES

OS Artificial Sequence
PN JP 199341989-A/65
PD 14-DEC-1999
PF 16-MAR-1999 JP 1999069694
PR PI KOICHI INOUE
PC C12N15/09,C12M1/00,C12Q1/68,C12N15/00
CC
FH Key
Location/Qualifiers
FT source

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//cganism='Artificial Sequence'.
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OS Unidentified
PN JP 2000270867-A/65
PD 03-OCT-2000
PF 19-MAR-1999 JP 1999
PR KOICHI INOUE
PC C12NL5/09,CL2M1/00,CC Strandedness: Singlific CC Topology: Linear;
FH Key Local FT Source 1.
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synthetic construct
artificial sequences.
1 (bases 1 to 12)
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E64117
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Similarity 90.9%;
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C12N15/09,C12M1/00,C12Q1/68,C12N15/00
Strandedness: Single;
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                                                                                                                                                     /organism='Artificial
Location/Qualifiers
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/mol_type="genomic DNA"
/db_xref="taxon:32644"
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Location/Qualifiers
                                                                  /organism="synthetic construct"
/mol_type="genomic DNA"
/db_xref="taxon:32630"
  31.3%;
90.9%;
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Pred. No. 41;
  Score 9.4; Di Pred. No. 41;
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BD101930/c
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BD061483/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Method for discriminating microorganisms, apparatus for discriminating microorganisms, method for preparing data base for discriminating microorganisms, and recording medium recorded with program for discriminating microorganisms

AL Patent: JP 2001275700-A 10 09-OCT-2001;
SANYO ELECTRIC CO LTD, SOCIETY FOR TECHNO-INNOVATION OF AGRICULTURE FORESTRY AND FISHERIES

OS Artificial Sequence
JP 2001275700-A/10
PD 09-OCT-2001
PF 31-MAR-2000 JP 2000099482
PF 31-MAR-2000 JP 2000099482
PC C1201/68,C12M1/00,C12M1/34,C12N15/09,G06F17/30,C12N15/00 CC
Primer Location/Qualifiers
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BD061483
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   discriminating microorganisms, microorganisms program and record medium for recording the sa Patent: WO 0175156-A 10 11-OCT-2001; SANYO ELECTRIC CO LTD, SOCIETY FOR TECHNO INNO
                                                   Method of discriminating microorganisms, apparatus for discriminating microorganisms, method of making database for discriminating microorganisms, microorganisms discriminating
                                                                                                                      synthetic construct
synthetic construct
artificial sequences.
1 (bases 1 to 12)
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WO 0175156-A/10.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   /mol_type="genom.../
/db_xref="taxon:32630"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    /organism="synthetic construct"
/mol_type="genomic DNA"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Location/Qualifiers
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90.9%;
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   SOCIETY FOR TECHNO INNOVATION OF AGRICULTURE
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AUTHORS
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VERSION
KEYWORDS
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BD240229/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                          Mammalia; Eutheria; Primates; Catarrhini; H
1 (bases 1 to 10)
Roberts, B.L. and Shankara, S.
Preparation and use of superior vaccines
Patent: JP 2002534056-A 1647 15-OCT-2002;
GENZYME CORP
OS Homo sapiens (human)
PN JP 2002534056-A/1647
PD 15-OCT-2002
PF 18-JUN-1999 JP 2000554749
PF 18-JUN-1999 JP 2000554749
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OS Artificial Sequence
PN W0 0175156-A/10
PD 11-0CT-2001
PF 27-MAR-2001 W0 2001JP002516
PF 27-MAR-2000 JP 00P 099482
PI TAKAKAZU INOUE
PC C12Q1/68,C12N15/10,G01N33/48,G01N27,
Primer
                                                                                                                                                                     19-JUN-1998 US
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19-JUN-1998
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                                                                                                      G01N37/00
                                                                                                                                       C12N1/19
                                                                                                                      C12N1/21, C12N5/10, G01N33/15, G01N33/50, G01N33/53, G01N33/566,
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W 0175156-A/10
11-OCT-2001
27-MAR-2001 WO 2001JP002516
31-MAR-2000 JP 00P 099482
TAKAKAZU INOUE
C12Q1/68,C12N15/10,G01N33/48,G01N27/447,G06F17/30,C12M1/00
                                                      Key
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                                      source
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/organism='Homo sapiens
Location/Qualifiers
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/mol_type="genomic DNA"
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                                                                                                                                                                         ROBERTS, SRINIVAS SHANKARA
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90.9%;
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US 60/0900
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60/090078,19-JUN-1998
60/090076,19-JUN-1998
60/111715
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60/090080,19-JUN-1998
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60/090035,19-JUN-1998
60/089992,19-JUN-1998
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60/090041,19-JUN-1998 US
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                                                                        use of
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                                                   use of superior vaccines Location/Qualifiers
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Catarrhini; Hominidae;
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60/090040

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60/089853

60/090079 60/089993 60/090072 60/089991 60/090048 60/090043

60/089844 60/089833

(human)'

PC PC

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REFERENCE
AUTHORS
TITLE
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BD248497
                                         SOURCE
ORGANISM
                                                                                  ACCESSION
                                                                                                                RESULT 54
AR303300/c
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Best Local
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            AUTHORS
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T cells
1 (bases 1 to 10)
Shimamoto, A., Furuichi, Y.,
Watahiki, M.
                                                                                  AR303300
Sequence 25 from patent US
AR303300
                                                                                                                                                           N
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synthetic construct
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                                         Unknown
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                               Unclassified.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           rtificial sequences.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                           cells specific for target antigens and
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                                                                                                                                                           CAGACTTCA 10
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A61K39/395,A61P31/04,A61P31/10,A61P31/12,A61P35/00,C12N5/10,
                                                                                                                                                                                                                                                                                                                                                                               10-NOV-1998 JP 2000581183
MAURICE ZAUDERER
C12N15/09,A01K67/027,A61K35/76,A61K39/00,A61K39/04,A61K39/12,
                                                                                                                                                                                                    Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                /organism="Homo sapiens"
/mol_type="genomic DNA"
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                                                                                                                                                                                                                                                                                  /organism='Artificial Sequence'
                                                                                                                                                                                                                                          /organism="synthetic construct"
/mol_type="genomic DNA"
/db_xref="taxon:32630"
                                                                          GI:31692076
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100.0%; Pr
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Pred. No. 41;
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            Shibata,Y.,
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o. 41;
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            Funaki, H.,
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            Ohara, E.
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AUTHORS
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AX510716
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KEYWORDS
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JOURNAL
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TITLE
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Best Local
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Matches
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AX471386
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Sequence 4 from Patent WO0227027.
AX510716
                                                                                                                                                         Homo sapiens
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
                                                                                                                                                                                                                                                                                                                       N
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Method of screening for therapeutics Patent: WO 0227027-A 4 04-APR-2002; THE UNIVERSITY OF ROCHESTER (US)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             synthetic construct
synthetic construct
artificial sequences.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Method for synthesizing cDNA from mF
Patent: US 6544736-A 25 08-APR-2003;
                                                                                                        HENKEL KGAA (DE)
                                                                                                                 Hofmann,K., Conradt,M. and Petersohn,D Method for determining skin stress or Patent: WO 02053773-A 963 11-JUL-2002;
                                                                                                                                                                                                     Homo sapiens (human)
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Similarity 9; Conserv
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Conservative 0;
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963 from Patent WO02053773.
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/mol_type="genomic |
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/mol_type="unassigned DNA"
/db_xref="taxon:9606"
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/db_xref="taxon:32630"
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Sequence 4026 from Patent
AX626985
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae;
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/db_xref="taxon:9606"
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Sequence 9694 from Patent
AX632652
            Petersohn,D., Conradt,M. and Hofmann,K. Method for determining homeostasis of the skin Patent: WO 02053774-A 9694 11-UUL-2002; Henkel Kommanditgesellschaft auf Aktien (DE)
                                                                                  Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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/mol_type="unassigned DN
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Amerson,H.V., Wilcox,P., Sederoff,R.R.,
O'Malley,D.M. and Grattapaglia,D.
Methods for within family selection of d
perennials using genetic markers
Patent: US 5908978-A 20 01-JUN-1999;
                                                                                                                  1 (bases 1 to 10)
1 (bases 1 to 10)
1 (bases 1 to 10)
Reinath, A.P., Somai, B.M. and Dean, R.A.
Method of diagnosing gummy stem blight
chain reaction assay
Patent: US 6258537-A 6 10-JUL-2001;
Location/Qualifiers
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Unclassified.
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                     GAAGCCAGAC 1497
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GATGCCAGAC 10
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20 from patent US 5908978.
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/mol_type="unassigned DNA"
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                                                                                                                                                                                                                                                                                                                                                                                                                    /organism="unknown"
/mol_type="unassigned
                                                                                                                                                                                                                                                                                                                                                                                                                                                      Location/Qualifiers
                                                                                      /mol_type="unassigned DNA"
                                                                                                 organism="unknown"
                                                                                                                                                                                                                                        GI:16228965
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90.0%;
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                                                                 Length 10;
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BD240233
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Roberts, B.L. and Shankara, S. Preparation and use of superior vaccines patent: JP 2002534056-A 1651 15-OCT-2002; GENZYME CORP 2002534056-A/1651 15-OCT-2002; JP 2002534056-A/1651 PD 15-OCT-2002 PF 18-JUN-1998 US 60/09003, J9-JUN-1998 US 19-JUN-1998 US 60/090041, J9-JUN-1998 US 60/090035, J9-JUN-1998 US 19-JUN-1998 US 60/090035, J9-JUN-1998 US 19-JUN-1998 US 60/090035, J9-JUN-1998 US 19-JUN-1998 US 60/09000, J9-JUN-1998 US 19-JUN-1998 US 60/09000, J9-JUN-1998 US 19-JUN-1998 US 60/09000, J9-JUN-1998 US 19-JUN-1998 US 60/090042, J9-JUN-1998 US 19-JUN-1998 US 60/090044, J9-JUN-1998 US 19-JUN-1998 US 60/090044, J9-JUN-1998 US 19-JUN-1998 US 60/090044, J9-JUN-1998 US 19-JUN-1998 US 60/090076, J9-JUN-1998 US 19-JUN-1998 US 60/090078 19-JUN-1998 US 19-JUN-1998 US 60/090078 19-JUN-1998 US 19-JUN-1998 US 60/090078 19-JUN-1998 US 19-J
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                                                                                                                                E54684.1 GI:22556167
JP 2001211883-A/36.
Homo sapiens (human)
Homo sapiens
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae;
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JP 2002534056-A/1651.
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   Matsushima, K.,
                        Eukaryota; Metazoa; Chordata;
Mammalia; Eutheria; Primates;
1 (bases 1 to 10)
                                                                                                                                                                                                                                                                 E54684
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C12N1/21,C12N5/10,G01N33/15,G01N33/50,G01N33/53,G01N33/566,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          GAAGCCAGCC 10
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Preparation and use of superior vaccines Key Location/Qualifiers
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                                                                                                                                                                                                                                                                                                   normal liver
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98 US 60/990039,19-JUN-1998 US 60/99041,19-JUN-1998 US 60/99097,19-JUN-1998 US 60/089997,19-JUN-1998 US 60/089992,19-JUN-1998 US 60/089992,19-JUN-1998 US 60/089991,19-JUN-1998 US 60/089090,19-JUN-1998 US 60/090042,19-JUN-1998 US 60/090044,19-JUN-1998 US 60/090044,19-JUN-1998 US 60/090044,19-JUN-1998 US 60/090044,19-JUN-1998 US 60/090044,19-JUN-1998 US 60/090078,19-JUN-1998 US 60/090078,19-JUN-19-JUN-19-JUN-19-JUN-19-JUN-19-JUN-19-JUN-19-JUN-19-JUN-19
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   Hashimoto,S.,
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Kaneko, S.
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                                                              Craniata; Vertebrata; l
Catarrhini; Hominidae;
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Yamashita,T
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60/08993 PR
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60/08991 PR
60/08991 PR
60/090043 PR
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60/089844 PR
60/089833 PR
60/080077 PR
60/090045 PR
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ORGANISM
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ACCESSION
VERSION
                                                                                                                                                                            RESULT 67
AR303303
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Best Local (
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YAMASHITA
PC C12NI
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OS Homo sapiens (human)
N J 2001211883-A/36
DD 07-AUG-2001 JP 2000023170
PF 31-JAN-2000 JP 2000023170
                    Method for synthesizing cDNA from mRNA sample Patent: US 6544736-A 28 08-APR-2003; Location/Qualifiers
                                                             1 (bases 1 to 10)
Shimamoto, A., Furuichi, Y., Shibata, Y.,
                                                                                                                                            AR303303
Sequence 28 from patent US
AR303303
                                                                                                                                                                                                                                                                 28.0%;
Similarity 90.0%;
9; Conservation
                                                                                                                                                                                                                                                                                                                                             MO (bases 1 to 10)
MO (months) Mosquito and/or flea control
Patent: US 6337071-A 12 08-JAN-2002;
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 12 from patent AR181983
                                                                                                Unknown
                                                                                                                                 AR303303.1 GI:31692079
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                                                                                      Unclassified.
                                                                                                            Unknown.
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9; Conserv
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31-JAN-2000 JP 2000023170
KOJI MATSUSHIMA,SHINICHI HASHIMOTO,SHUICHI KANEKO,TARO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Conservative
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/mol_type="genomic DNA"
/db_xref="taxon:9606"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Location/Qualifiers
 /organism="unknown'
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/mol_type="unassigned DNA"
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                                                                                                                                                                                                                                                                                       Score 8.4; DB 1;
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                                                                Funaki,H.,
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                                                                  Ohara, E.
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           RESULT 70
AR382219
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1486 AAGAAGCCAG 1495
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Bullerdiek, Van de Ven, W.J.M., Schoenmakers, H.F.P.M. and Mols, R. Multiple-tumor aberrant growth genes
Patent: US 6544784-A 110 08-APR-2003;
                                                                                                                                                                                                                                                                                                                                   Sequence 110 from patent AR304485
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Shimamoto, A., Furuichi, Y., Shibata, Y.,
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Similarity 90.0%;
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/mol_type="genomic |
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        /mol_type="genomic DNA"
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Pred. No. 51;
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              PAT 18-DEC-2003
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AX152377
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Sequence 141 from Patent WO0138577.
AXI52226
AXX152226.1 GI:14533877
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Keinath, A.P., Somai, B.M. and Dean, R.A.

Method of diagnosing gummy stem blight

chain reaction assay

Patent: US 6610487-A 6 26-AUG-2003;

Location/Qualifiers
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         Human transcriptomes
Patent: WO 0138577-A 292 31-MAY-2001;
The Johns Hopkins University (US)
Location/Qualifiers
                                                                                         Eukaryota; Metazoa; Chordata; Mammalia; Eutheria; Primates;
                                                                                                                  Homo sapiens (human)
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Sequence 292 from Patent
AX152377
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Patent: WO 0138577-A 141 31-MAY-2001;
The Johns Hopkins University (US)
Location/Qualifiers
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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                                                              Velculescu, V.E., Vogelstein, B. and Kinzler, K.W
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/mol_type="unassigned DNA"
/db_xref="taxon:9606"
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AX153162/c
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Eukaryota; Metazoa; Chordata;
Mammalia; Eutheria; Primates;
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Sequence 1077 from Patent W00138577,
AX153162
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Patent: WO 0138577-A 1077 31-MAY-2001;
The Johns Hopkins University (US)
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                                              Similarity 9; Conserv
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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/db_xref="taxon:9606"
                                                                                                /organism="Homo sapiens"
/mol_type="unassigned DN
/db_xref="taxon:9606"
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Velculescu, V.B., Vogelstein, B. and Kinzler, K.W. Characterization of the yeast transcriptome Patent JP 2001599017-A 53 10-JUL-2001;

THE JOHNS HOPKINS UNIVERSITY SCHOOL OF MEDICINE OS Saccharomyces cerevisiae (yeast)

PN JP 2001509017-A/53

PD 10-JUL-2001

PF 22-JAN-1998 JP 1998532117
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D 10-UUL-2001

PF 22-JAN-1998 JP 1998532117

PF 23-JAN-1997 US 60/035917

PR 23-JAN-1997 US 60/035917

PR 23-JAN-1997 US 60/035917

PI VICTOR E VELCULESCU, BERT VOGELSTBIN, KENNETH W KINZLER PC C12N15/1, C12N15/31, C07K14/395, C12Q1/68, C12Q1/02 CC C12N15/10, C12N15/31, C07K14/395, C12Q1/68, C12Q1/02 CC Characterization of the yeast transcriptome Location/Qualifiers

Location/Qualifiers
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Saccharomyces cerevisiae
Eukaryota; Fungi; Ascomycota; Saccharomycotina; Saccharomycetes;
Saccharomycetales; Saccharomycetaceae; Saccharomyces.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    BD065117
Characterization of the yeast transcriptome.
BD065117
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Sequence 63 from Patent WO0212498.
AX377325
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                                                           Similarity 9; Conserva
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                                                                                                                                                                                    /organism='Saccharomyces
Location/Qualifiers
                                                                                                                      /organism="Saccharomyces
/mol_type="genomic DNA"
/db_xref="taxon:4932"
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/mol_type="unassigned DNA"
/db_xref="taxon:9606"
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Pred. No. 51;
                                                            Mismatches
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BD167088/c
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PC C12N;
PC C12N;
PC C12N;
PC Human
FH Key
FT Source
                                                                                                                                                       VECTOR FOR THE EXPRESSION IN YEASTS OF INTERLEUKINE-2, YEASTS AND METHOD FOR PREPARING INTERLEUKINE-2 Patent: WO 8503723-A 10 29-AUG-1985;
Location/Qualifiers
                                                                                                                                                                                                                                                                                           Nucleotide sequence 10
A02163
A02163.1 GI:410850
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OS Homo sapiens (human)
PN JP 2002209591-A/633
PD 30-JUL-2002
PF 19-JAN-2001 JP 2001012328
PF 19-JAN-2001 JP 2001012328
PI KOJI MATSUSHIMA, SHINICHI HASHIMOTO, SHUICHI KANEKO, TARO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              BD167088
BD167088.1 GI:27872900
JP 2002209591-A/633.
                                                                                                                                                                                                                                        unclassified.
                                                                                                                                                                                                                                                   unidentified
                                                                                                                                                                                                                                                                   unidentified
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                                                   Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          /organism="unidentified"
/mol_type="genomic DNA"
/db_xref="taxon:32644"
                                                                                                     /organism="unidentified"
/mol_type="unassigned DN
/db_xref="taxon:32644"
                                                              28.0%;
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O; Mismatches
                                                  Score 8.4; DB Pred. No. 54; 0; Mismatches
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Location/Qualifiers
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Sequence
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Lemoine, Y., Sondermeyer, P., Loison, G., Aigle, M. and Lecocq, J.P.
Yeast-expression vectors for interleukin-2, transformed yeasts and
process for the preparation of interleukin-2
Patent: EP 0152358-A 9 21-AUG-1985;
                                                                                                                                                                                                                                                                                              1 (bases 1 to 11)
Wang,C.-G. and Hepburn,A.G.
Genetic sequence assay using DNA triple strand formation
Patent: US 5861244-A 342 19-JAN-1999;
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               unidentified unclassified
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                                 Unknown
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AR353840
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 1 (bases 1 to Baric, R.S. and
                        Unclassified.
                                                Unknown.
                                                                      AR353840.1 GI:33759907
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               /organism="unidentified"
/mol_type="unassigned DNA"
/db_xref="taxon:32644"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Location/Qualifiers
                                                                                                                                                                                                                                                            /mol_type="unassigned DNA"
                                                                                                                                                                                                                                                                         organism="unknown"
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Yount, B.
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                                                                                                                                                                                              Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae;
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                                                            /organism="Homo sapiens"
/mol_type="unassigned DNA"
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/mol_type="genomic |
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/db_xref="taxon:9606"
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AX471384/c
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Sequence 16 from Patent EP1262565.
AX616455
AX616455.1 GI:28447498
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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Sequence 961 from Patent WO02053773.
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae;
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            1632
                                                                                                                                                                                              /organism="Homo sapiens"
/mol_type="unassigned DNA"
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/db_xref="taxon:9606"
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Homo sapiens
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Sequence 3005 from Patent
AX625964
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Method for determining homeostasis of the skin
Patent: WO 02053774-A 2530 11-JUL-2002;
Henkel Kommanditgesellschaft auf Aktien (DE)
Location/Qualifiers
              Petersohn,D., Conradt,M. and Hofmann,K. Method for determining homeostasis of the skin Patent: WO 02053774-A 3005 11-JUL-2002;
                                                                                                                                                                                                                                                                                                                                   28.0%;
Similarity 90.0%;
9; Conservation
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                                                                          Eukaryota; Metazoa; Chordata;
Mammalia; Eutheria; Primates;
                                                                                       Eukaryota; Metazoa;
                                                                                                                                                    AX625964.1 GI:28454002
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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Location/Qualifiers
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
Henkel Kommanditgesellschaft auf Aktien
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AX626847/c
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AX627093/c
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Homo sapiens
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Patent: WO 02053774-A 4134 11-JUL-2002;
Henkel Kommanditgesellschaft auf Aktien (DE)
Location/Qualifiers
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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/db_xref="taxon:9606"
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/mol_type="unassigned DNA"
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Sequence 4328 from Patent WO02053774.
AX627287
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae;
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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Sequence 4967 from Patent
AX627926
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             Petersohn,D., Conradt,M. and Hofmann,K. Method for determining homeostasis of the skin Patent: WO 02053774-A 5549 11-JUL-2002; Henkel Kommanditgesellschaft auf Aktien (DE) Location/Qualifiers
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae;
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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/mol_type="unassigned DNA"
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae;
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Mammalia; Eutheria; Primates;
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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                                            Petersohn,D., Conradt,M. and Hofmann,K. Method for determining homeostasis of the skin Patent: WO 02053774-A 9054 11-JUL-2002; Henkel Kommarditgesellschaft auf Aktien (DE) Location/Qualifiers
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae;
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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/note= "U-rich
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HIV-2 (see
The invention relates to a method of determining if a compound binds to a target RNA by treating a test compound with a reporter (R) labelled with a donor or acceptor group and labelled target RNA, labelled with the complementary donor or acceptor group, and measuring the fluorescence from fluorescent groups associated with a compound:target RNA complex in presence of the test compound and comparing the result with a standard.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           New HIV growth inhibiting oligo:nucleotide(s) - comprisin sequences capable of binding to tat protein within cells, to identify cpds. with tat binding.
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02-MAR-1999;
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11-APR-2000
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                                                                                                            ing compounds that bind infections, tumors and
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nilarity 76.5%;
Conservative
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Best Local Similarity
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30-OCT-2000;
27-MAR-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Chromosomal genomic alteration; genome altering oligonucleotide; PCR; ss o-methyl modification; LNA modification; phosphorothioate linkage; DNA repair; DNA alteration; environmental tolerance; hygromycin; primer amino acid over production; herbicide resistance; glyphosate resistance; imidazolinone herbicide resistance; sulphonylurea herbicide resistance; porphyric herbicide resistance; sulphonylurea herbicide resistance; porphyric herbicide resistance; prophyric herbicide resistance; production; modified oil production; modified starch production; modified floral morphology; male-sterile plant; albino mutant; altered floral morphology; male-sterile plant; albino mutant; increased stearate production; reduced plantiate production; albino plant; altereated production; production; reduced linolenic acid production;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         The oligonuclectides AAZ59070-Z59071 anneal to form a double stranded oligonuclectide containing the HIV-1 trans-activation regulatory region (TAR) to which the HIV-1 Tat protein binds. The complex is labelled with 6-carboxyfluorescein and is used as a target for the binding of a labelled ADP-1 protein. Detection of the complex is by fluorescence resonance energy transfer (FRET). The method is used to identify compounds that interfere with interaction between the target RNA and ligands or proteins. Compounds that are identified are potentially useful for treating infections (viral, bacterial or fungal), cancer and autoimmune diseases. The compounds are preferably directed to the TAR and RRE regions of human immunodeficiency virus RNA and inhibit viral replication. (Updated on 15-SEP-2003 to standardise OS field)
                                                                                                                  New oligonucleotides with moderneating plants with desired nutritional value, herbicide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     09-APR-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 17 BP; 5
                                                                                            production.
                                                                                                                                                                                                                                           WPI; 2002-106307/14.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       01-JUN-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WO200192512-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Solanum tuberosum.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                photosynthetic process.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ABK26439 standard; DNA; 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       1490
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                                                                                                                                                                                                                                                                                                        EB,
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                                                                                                                                                                                                                                                                                                     Gamper HB,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  production genome altering oligonucleotide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (first entry)
                                                                                                                                                                                                                                                                                                                                                                     DELAWARE
                                                                                                                                                                                                                                                                                                                                                                                                                           2000US-0208538P.
2000US-0244989P.
2001US-00818875.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       2001WO-US017672.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             46.0%;
                                                                                                                                                                                                                                                                                                           Rice
                                                                                                                     modified nuclease-resistant termini, red phenotypes, e.g. stress tolerance, ide or disease resistance, or modified
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           BP.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                <u>ب</u>
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               hygromycin; primer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        #95
                                                                                                                            modified
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                                                                                                            improved doil
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Kmiec EB,

Gamper HB,

Rice MC,

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ABK26440/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Consist of o-methyl modification, an LNA modification. The phosphorochioate linkages on a terminus, or a combination of these modifications of the oligonucleotide and further comprises consist of o-methyl modification, an LNA modification, two or more compiss of these modifications. The oligonucleotide. The chemical modifications of the oligonucleotide are useful for consist of o-methyl modification, an LNA modification, two or more composite of these modifications. The oligonucleotides are useful for content of these modifications. The oligonucleotides are useful for creating plants with desired coligonucleotides are particularly useful for creating plants with desired content of plants or conferring amino acid over production), herbicide resistance (e.g. conferring amino acid over production), herbicide resistance (e.g. glybhosate resistance, imidazolinone and sulphonylurea herbicide resistance, porphyric herbicide resistance or triazine resistance), constants or production modified starch production (e.g. increased starch or production of modified starch production (e.g. reduced palmitate, increased stearate or reduced linolenic acid). The oligonucleotides are also useful for producing albino mutants for the altering oligonucleotide production of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           RESULT 4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Matches
                                                                                                                       01-JUN-2000;
30-OCT-2000;
27-MAR-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       altered floral morphology; male-sterile plant; albino mutant; modified fatty acid content; reduced palmitate production; albino increased stearate production; reduced linolenic acid production;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              o-methyl modification; INA modification; phosphorothioate linkage;
DNA repair; DNA alteration; environmental tolerance; hygromycin-B;
abiotic stress tolerance; improved nutritional value; hygromycin; primer
amino acid over production; herbicide resistance; glyphosate resistance;
imidazolinone herbicide resistance; sulphonylurea herbicide resistance;
porphyric herbicide resistance; triazine resistance; disease resistance;
modified oil production; modified starch production; waxy starch;
                                                                                                                                                                                                                                                         01-JUN-2001;
                                                                                                                                                                                                                                                                                                                        06-DEC-2001.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Solanum tuberosum.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Chromosomal genomic alteration;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Waxy starch production genome altering oligonucleotide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ABK26440
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         The invention relates to an oligonucleotide for targeted alteration
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         photosynthetic process.
                                                               (UYDE ) UNIV DELAWARE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       1485 CAAGAAGCCAGACTTCA 1501
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       15;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            standard; DNA; 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CAAGAAGCTAAACTTCA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          oligonucleotide of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              BP; 8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Conservative
                                                                                                                          2000US-0208538P.
2000US-0244989P.
2001US-00818875.
                                                                                                                                                                                                                                                         2001WO-US017672
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              46.0%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       <u>.</u>.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Pred
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Score 13.8;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         genome altering oligonucleotide;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        No.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            albino plant;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PCR; ss;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         primer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               of a
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2002-106307/14.

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RESULT 5
AAZ59072
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   genetic sequence, which comprises a single-stranded oligonucleotide having a DNA domain. The DNA domain has at least one mismatch with respect to the genetic sequence to be altered and further comprises chemical modifications of the oligonucleotide. The chemical modifications consist of o-methyl modification, an LNA modification, two or more phosphorothioate linkages on a terminus, or a combination of any two or more of these modifications. The oligonucleotides are useful for directing repair or alteration of plant genetic information. The oligonucleotides are are particularly useful for creating plants with desired phenotypes, e.g. environmental or abiotic stress tolerance, improved nutritional value (e.g. altering amino acid content of plants or conferring amino acid over production), herbicide resistance (e.g. glyphosate resistance, imidazolinone and sulphomylurea herbicide resistance, porphyric herbicide resistance or triazone resistance, and the production of triazone resistance.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               disease resistance, modified oil production, modified starch production (e.g. increased starch or production of waxy starch), altered floral morphology (e.g. male-sterile plants) or modified fatty acid content (e.g. reduced palmitate, increased stearate or reduced linolenic acid). The oligonucleotides are also useful for producing albino mutants for the analysis of photosynthetic processes. This sequence represents a genome altering oligonucleotide of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                New oligonucleotides with modified nuclease-resistant termini, useful for creating plants with desired phenotypes, e.g. stress tolerance, improved nutritional value, herbicide or disease resistance, or modified oil production.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               The invention relates to an oligonucleotide for targeted alteration of genetic sequence, which comprises a single-stranded oligonucleotide
                                                                      05-JUN-1998;
02-MAR-1999;
                                                                                                                                                                                                                                                                 Antiviral; antibacterial; antifungal; anticancer; detection; TAR; fluorescence resonance energy transfer; tat; HIV-1; Rev response e autoimmune disease; trans-activation regulatory region; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 17
 Karn J,
                                                                                                                          04-JUN-1999;
                                                                                                                                                               16-DEC-1999.
                                                                                                                                                                                                WO9964625-A2
                                                                                                                                                                                                                                 Human immunodeficiency virus 1.
                                                                                                                                                                                                                                                                                                                                       HIV-1 TAR oligonucleotide target sequence
                                                                                                                                                                                                                                                                                                                                                                              15-SEP-2003
11-APR-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                  AAZ59072;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAZ59072 standard;
                                     (RIBO-) RIBOTARGETS LTD
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1485 CAAGAAGCCAGACTTCA 1501
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               7; Page 151; 220pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CAAGAAGCTAAACTTCA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   BP;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Conservative
                                                                                                                                                                                                                                                                                                                                                                            (revised)
(first entry)
                                                                    98GB-00012196.
99GB-00004790.
                                                                                                                          99WO-GB001761
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 A;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 RNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             46.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 2 C; 4 G; 8
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Score 13.8;
Pred. No. 9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 T; 0 U;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0 Other;
                                                                                                                                                                                                                                                                                                                                           #3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             <u>ب</u>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Length 17;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Indels
                                                                                                                                                                                                                                                                    Rev response element; ion; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Gaps
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Identifying compounds that bind to target RNA, potentially useful
                                                                                infections, tumors and
                                                                                 autoimmune diseases.
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Example; Page 31; 82pp; English.

CC target RNA by treating a test compound with a reporter (R) labelled with the CC a donor or acceptor group and labelled target RNA, labelled with the CC complementary donor or acceptor group, and measuring the fluorescence CC from fluorescent groups associated with a compound: the RNA complex in CC presence of the test compound and comparing the result with a standard. CC The oligonucleotides AAZ59070-Z59071 annual to form a double stranded CC oligonucleotide containing the HIV-1 trans-activation regulatory region CC (TAR) to which the HIV-1 Tat protein binds. The complex is labelled with CC (arboxyfluorescein and is used as a target for the binding of a CC labelled ADP-1 protein. Detection of the complex is by fluorescence CC resonance energy transfer (FRET). The method is used to identify compounds that interfere with interaction between the target RNA and CC ligands or proteins. Compounds that are identified are potentially useful CC autoimmune diseases. The compounds are preferably directed to the TAR and CC RETIONS of Auman immunodeficiency virus RNA and inhibit viral CR replication (Indated on 15-SEP-2007 for strangardian of finishit viral). The invention relates to replication. (Updated on 15-SEP-2003 to standardise a method of determining if a compound binds to porter (R) labelled with SO

Sequence 18 BP; 5 A; 4 C; 6 G; 0 T; 3 U; 0 Other;

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Best Local Similarity
Matches 13; Conserv
                                                          Query Match
                   1490 AGCCAGACTTCAGCAGC 1506
AGCCAGAUJUGAGCAGC 17
                                       Conservative
                                               46.0%;
76.5%;
                                      2; Mismatches
                                                 Score 13.8;
Pred. No. 9
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                                                           DB
                                                          1;
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                                                          Length 18
                                        Indels
                                       0;
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RESULT 6
AAF74454
ID AAF7
XX
AAF74454
AC AAF7
XX
AC AAF7
XX
DT 09-1
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DT 09-1
XX
DT U9-1
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PD 1:
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PD 1:
X
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       09-MAY-2001
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Human; PRO; PROX; cytostatic; immunomodulatory; reproduction; gene therapy; cell proliferation; differentiation disorder; c immune associated disorder; gestational disease; pre-clampsia PCR primer; sequencing primer; ss. pre-clampsia;

cancer;

Homo sapiens

WO200110902-A2

15-FEB-2001.

11-AUG-2000; 2000WO-US021857.

11-AUG-1999; 10-AUG-2000; 2000US-00635949 99US-0148433P

CURAGEN CORP.

Shimkets RA, Fernandes

WPI; 2001-147509/15.

Nucleic acids encoding secreted polypeptides, polypeptides, useful for treating a syndrome associated disorder, e.g. cancer. , designated PROX associated with PROX Ø PROX-

Example

2

Page 119; 166pp;

English

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AAF74457/c

ID AAF74457 standard; D

XX

AC AAF74457;

XX

DT 09-MAY-2001 (first

XX

DT 09-MAY-2001 (first

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DT 09-MAY-2001 (first

XX

DE Human; PRO; PROX; cy

KW Human; PRO; PROX; cy

KW Gene therapy; cell p

KW PCR primer; sequenci

XX

Homo sapiens.

XX

PN W0200110902-A2.

XX

PT 1-AUG-1909; 99US-

PR 11-AUG-1909; 99US-

PR 11-AUG-2000; 2000W0-

XX

PN W0200110902-A2.

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PN W0
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Best Local
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                                                                                                                                                                                                                                                          Nucleic acids encoding secreted polypeptides, designated PROX polypeptides, useful for treating a syndrome associated with associated disorder, e.g. cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  gene therapy; cell proliferation; differentiation disorder; ca immune associated disorder; gestational disease; pre-clampsia;
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gene therapy; cell proliferation; differentiation disorder; c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Human PRO2 gene-specific sequencing primer SEQ ID NO:43
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                                    The present invention describes isolated nucleic acids encoding secreted polypeptides, designated PROX perpeptides (i.e. a PRO polypeptide where X is an integer from 1 to 17). PROX polypeptides have cytostatic, immunomodulatory and reproduction activities, and can be used in gene
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             11-AUG-1999; 99US-0148433P.
10-AUG-2000; 2000US-00635949.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 primer; sequencing
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                                                                                                                                                                                               Page 119; 166pp; English.
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      PROX antagonists and PROX agonists.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           DNA;
                                                                                                                                                                                                                                                       e.g.
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88.2%;
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No. 9.
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                                                                                                                                                                                                                                                                                              associated with a
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   an be used in gene
PROX polypeptides,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      cancer;
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GGAGCCAGACTTCA

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RESULT 8
AAX577566
ID AAX577566
ID AAX577566
ID AAX577566
ID AAX5
XX AAX5
XX AAX5
XX AAT5
XX AAT5
XX AAX5
XX AAY5
XX AAX5
XX A
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Best Local S
Matches 15
                                                                Query Match
Best Local Similarity
Matches 13; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                nucleic acids and antibodies are useful in the manufacture of a medicament for treating a syndrome associated with a PROX-associated disorder, e.g. a cell proliferation and/or differentiation disorder (e.g. famune associated disorders) and a gestational disease (e.g. pre-clampsia). They are also used for screening for a modulator of activity or of latency or predisposition to a PROX-associated disorder AAF74432 to AAF74448 encode the specifically claimed human PROX polypeptides PRO1 to PRO17 given in AAB70531 to AAB70547. The present sequence represents a primer used in an example from the present envent.
                                                                                                                                                                                                                            Sequences AAX57562-X57571 represent antisense oligonucleotides targeted to a region spanning 4-9 codons downstream of the AUG translation initiation codon of the human insulin-like growth factor-1 receptor (IGF-IR) gene. The antisense oligonucleotides inhibit the expression of IGF-IR, which in turn induces apoptosis, especially in a tumour cell. The oligonucleotides can be administered via a liposome
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AAX57566;
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                                                                                                                                                                       Sequence 15
                                                                                                                                                                                                                                                                                                                                                                                                                    Disclosure; Page 16; 23pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         oligonucleotides.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human insulin-like growth factor-1 receptor gene antisense
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             1488
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GAAGCCAGACTTCA 1501
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                                                                                                                                                                            BP; 4 A; 4 C; 4 G; 3 T; 0 U; 0 Other;
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                                                                     Conservative
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88.2%;
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                                                                                              Score 12.4;
Pred. No. 1
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Pred. No. 9.
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                                                                                                                       Length 15;
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                                                                                                                                     RESULT 9
AAF49086/c
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                                                                                                                                                                               The present invention relates to a method for ameliorating the effects of skin disorders. The method comprises contacting the skin with an antisense oligonucleotide, (for Insulin-like Growth Factor [IGF]-1 receptor, IGF binding protein [IGFBP]-2 or IGFBP3], which is capable of inhibiting or reducing growth factor mediated cell proliferation, inflammation and/or other disorders. The present sequence is an oligonucleotide which can be used to design the antisense oligonucleotides of the present invention (see AAP45151 and AAF45153-01igonucleotides of the present invention (see AAP45151) and AAF45153-174510]. The method is useful for ameliorating the effects of psoriasis, inthibyosis, pityriasis, ruba, pilaris, serborrhoea, keloids, keratosis, inthibyosis, pityriasis, ruba, pilaris, serborrhoea, keloids, keratosis, anyperneovascular condition such as a neovascular condition of the retina, brain or skin, growth factor-mediated malignancies, other sclerotic disease, kiney disease, hyperproliferation of the inside of blood
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Ameliorating the effects of a disorder, e.g. psoriasis, by administering UV (ultra-violet) treatment (optional) and an antisense nucleic acid that inhibits or reduces growth factor mediated cell proliferation and/or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Antisense therapy; antiproliferative; antiinflammatory; antipsoriatic; cytostatic; dermatological; cardiant; virucide; ophthalmological; keloid; skin disorder; Insulin-like Growth Factor 1 receptor; IGF-1; pityriasis; IGF binding protein; IGFBPB2; IGFBB3; inflammation; psoriasis; pilaris; growth factor mediated cell proliferation; ichthyosis; serborrhoea; ruba; keratosis; neoplasia; scleroderma; wart; skin cancer; sclerotic disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       keratosis; neoplasia; scleroderma; wart; skin cancer; sc. hyperneovascular condition; hyperplasia; kidney disease; neovascular condition of the retina; ss.
                                                                                                                                   Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                             Example
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Wraight CJ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     21-JUN-2000; 2000WO-AU000693.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                               inflammation.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       28-DEC-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Homo sapiens
                                                                                                                                                                     vessels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (MURD-)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              IGF-I oligonucleotide #46.
                                                                                 Local Similarity
                               1488
15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    MURDOCH CHILDRENS RES INST.
                                                                                                                                                                                                                                                                                                                                                                                                                                             8;
                                                                                                                                                                   or any
                               GAAGCCAGACTTCA 1501
                                                                                                                                     15
GGAGCCAGACTTCA 2
                                                                                                                                                                                                                                                                                                                                                                                                                                             Page 61; 201pp; English.
                                                                                                                                   B₽;
                                                                 Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Werther
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                                                                                                                                     G; 4 T;
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Pred. No. 1
                                                                   Mismatches
                                                                              No. 15;
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                                                                                                                                     0 Other;
                                                                                                 DB 1;
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                                                                                               Length 15;
                                                                   Indels
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                                                               Gaps
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RESULT 10 AAF49087/c

RESULT 11
AAZ64832/c
ID AAZ648
XX
AC AAZ648
XX

AAZ64832 standard; RNA; 14

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                                                                                                                                                                        skin disorders. The method comprises contacting the skin with an antisense oligonucleotide, (for Insulin-like Growth Factor [IGF] 1 receptor, IGF binding protein [IGFBD] -2 or IGFBD3), which is capable of inhibiting or reducing growth factor mediated cell proliferation, inflammation and/or other disorders. The present sequence is an oligonucleotide which can be used to design the antisense oligonucleotides of the present invention (see AAF45151 and AAF45153-F45161). The method is useful for ameliorating the effects of psoriasis, ichthyosis, pityriasis, ruba, pilaris, serborrhoea, keloids, keratosis, neoplasias, scleroderma, warts, benign growths, cancers of the skin, a hyperneovascular condition such as a neovascular condition of the retina, brain or skin, growth factor-mediated malignancies, other sclerotic disease, kidney disease, hyperproliferation of the inside of blood received the sease.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Antisense therapy; antiproliferative; antiinflammatory; antipsoriatic; cytostatic; dermatological; cardiant; virucide; ophthalmological; keloid; skin disorder; Insulin-like Growth Factor 1 receptor; IGF-1; pltyriasis; IGF binding protein; IGFBP-2; IGFBP3; inflammation; psoriasis; pilaris; growth factor mediated cell proliferation; ichthyosis; serborrhoea; ruba; keratosis; neoplasia; scleroderma; wart; skin cancer; sclerotic disease; hyperneovascular condition; hyperplasia; kidney disease; hyperneovascular condition of the retina; ss.
                                                                                                                                                               vessels o
                                                                                                                               Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Ameliorating the effects of a disorder, e.g. psoriasis, by administering UV (ultra-violet) treatment (optional) and an antisense nucleic acid that inhibits or reduces growth factor mediated cell proliferation and/or
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                                                                                                                                                                                                                                                                                                                                                                                                                                Example 8; Page 61; 201pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                21-JUN-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                 present
 14
                                                               Similarity
13; Conserv
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                                                                                                                                                                 ဝူ
                                GAAGCCAGACTTCA 1501
                                                                                                                                15
 GGAGCCAGACTTCA
                                                                                                                                                                 any other
                                                                                                                               BP;
                                                                                                                                                                                                                                                                                                                                                                                 invention relates to a method for ameliorating the effects ers. The method comprises contacting the skin with an
                                                                 Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Werther GA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (first
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                                                                                                                               2 A;
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                                                                             41.3%;
                                                                                                                               4 C;
                                                                                                                                                               hyperplasia
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                                                                                                                               G; 4 T;
                                                               <u>.</u>
                                                                             Pred.
                                                                                                Score 12.4;
                                                                   Mismatches
                                                                                    No.
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                                                                                                DB 1;
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                                                                                                Length 15;
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of.

28-MAR-2000

(first

entry)

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ABXO1669/c
ID ABXO166
XX ABXO16
XX ABXO16
XX ABXO16
XX ABXO16
XX BEZYMA
CR HEPATİ
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KW HCV ri
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                                                                                                                                                                                                                                                                                                                               Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                   The present sequence represents the preferred target sequence of an enzymatic nucleic acid, especially a hairpin ribozyme, which cleaves the Hepatitis C virus (HCV) RNA sequence at the base position given in the descriptor line. The HCV sequence was screened for optimal ribozyme target sites using a computer folding algorithm and regions of the mRNA which did not form secondary folding structures and contained potential ribozyme cleavage sites were identified. Ribozymes were synthesised to target these sites and their activities optimised by either varying the length of the binding arms or by modification to prevent degradation by nucleases. The ribozymes of the invention inhibit gene expression and/or viral replication, and are used to treat diseases associated with Hepatitis C virus (HCV) infection, e.g. cirrhosis, liver failure and hepatocellular carcinoma. The ribozymes may be used in combination with
          Enzymatic nucleic acid; RNA cleavage; Hepatitis C virus infection; HCV ribozyme; HCV expression; HCV replication; cirrhosis; virucide; liver failure; hepatocellular carcinoma; HCV infection; drug therap
                                                                                                                                                                ABX01669 standard;
                                                                                                                                                                                                                                                                                                                                                           Sequence 14 BP; 3 A; 2 C; 6 G; 0 T; 3 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Claim 2; Page 99; 123pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Novel ribozymes for the treatment of hepatitis C infection.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          04-NOV-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Enzymatic nucleic acid; hammerhead ribozyme; virus replication; cleavag cirrhosis; liver failure; hepatocellular carcinoma; interferon; cancer; autoimmune disease; ss.
                                                                                                      23-DEC-2002
                                                                                                                                    ABX01669;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   18-SEP-1998;
25-FEB-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Substrate
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                                                                                                                                                                                                                                                                     CTTCAGCAGCCA 1508
                                                                                                                                                                                                                                           CTTCAGCAGCCA 1
                                                                                                                                                                                                                                                                                                                                                                                         n to treat HCV infection, other infectious diseases, and cancer
                                                                         a
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           for hairpin ribozyme which cleaves HCV at nt.
                                                                         virus
                                                                                                                                                                                                                                                                                                   Conservative
                                                                                                     (first
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99US-00257608.
99US-00274553.
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                                                                         substrate
                                                                                                                                                                RNA;
 interferon alpha;
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                                                                         #154 for HCV hairpin
                                                                                                                                                                                                                                                                                                                Score 12; DB
Pred. No. 16;
                                                                                                                                                                                                                                                                                                   Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                diseases
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                                                                                                                                                                                                                                                                                                                               Length 14;
                                                                         ribozyme #154
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              therapy;
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Matches 12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   enzymatic nucleic acid or ribozyme is in a hammerhead (HH) or hairpin (HP) motif where the binding arms comprise sequences complementary to one of the substrate sequences defined in the specification. The HCV ribozymes are useful for modulating the expression and/or replication of HCV. They can be used to treat cirrhosis, liver failure and/or hepatocellular carcinoma. The HCV ribozymes are also useful for treating a condition associated with HCV infection in conjunction with one or more other drug therapies, particularly type I interferon, especially interferon alpha, beta or gamma or consensus interferon. The present sequence represents a substrate for a HCV hairpin (HP) ribozyme. Note: Some of the sequence data for this patent did not form part of the printed specification. The complete sequence data for this patent was obtained in electronic format directly from the USPTO web site at
IGF binding protein; IGFBP-2; IGFBP3; inflammation; psoriasis; pilar growth factor mediated cell proliferation; ichthyosis; serborrhoea; keratosis; neoplasia; scleroderma; wart; skin cancer; sclerotic dise
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               New ribozymes targeting RNA derived from hepatitis C virus inhibit viral replication and are useful to treat hepatitis C virus infections and cirrhosis, liver failure or hepatocellular carcinoma.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WPI;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           interferon gamma; consensus interferon; hepatotropic; antiinflammatory; substrate; hairpin ribozyme; HP ribozyme; ss.
                                               Antisense therapy; antiproliferative; antiinflammatory; antipsoriatic; cytostatic; dermatological; cardiant; virucide; ophthalmological; keloid; skin disorder; Insulin-like Growth Factor 1 receptor; IGF-1; pityriasis;
                                                                                                                                               30-MAR-2001
                                                                                                                                                                               AAF49084;
                                                                                                                                                                                                             AAF49084 standard; DNA; 15
                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Claim 2; Page 62; 80pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  The present invention relates to enzymatic nucleic acids which specifically cleave RNA derived from Hepatitis C virus (HCV).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Blatt L,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                27-JUN-2002
                                                                                                               IGF-I oligonucleotide #44.
                                                                                                                                                                                                                                                                                                                                                                                                                                                       seqdata.uspto.gov/psipsDIDEntry.html
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PAVCO P A.
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                                                                                                                                                                                                                                                                                                                                                         ilarity 100.0%;
Conservative (
                                                                                                                                                                                                                                                                                                                                                                                                                        BP; 3 A;
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                                                                                                                                                                                                                                                                                                                                                                                          40.0%;
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; Pred. No. 16;
0; Mismatches
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                pilaris;
noea; ruba;
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ARESULT 14
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XX AAF490
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XX 30-MAR
XX IGF-I
XX Antise
KW Cytost
KW Skin d
KW Skin d
KW Skin d
KW IGF b
KW Growth
KW RETATO
KW NPDETN
KW NPDETN
KW NEOVAS
XX OS HOMO 8
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CC The present invention relates to a method for ameliorating the effects of CC skin disorders. The method comprises contacting the skin with an CC antisense oligonuclectide, (for Insulin-like Growth Factor [IGF]-1 CC receptor, IGF binding protein [IGFBP]-2 or IGFBP3), which is capable of inhibiting or reducing growth factor mediated cell proliferation, CC inflammation and/or other disorders. The present sequence is an CC oligonuclectide which can be used to design the antisense CC oligonuclectide which can be used to design the antisense CC oligonuclectide with the present invention (see AAF45151 and AAF45153-CC [AF45161]. The method is useful for ameliorating the effects of psoriasis, CC inthyposis, pityriasis, ruba, pilaris, serborrhoea, keloids, keratosis, CC neoplasias, scleroderma, warts, benign growths, cancers of the skin, a CC hyperneovascular condition such as a neovascular condition of the retina, CC disease, kidney disease, hyperproliferation of the inside of blood cxx vessels or any other hyperplasia
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
Best Local
                                                       Antisense therapy; antiproliferative; antiinflammatory; antipsoriatic; cytostatic; dermatcological; cardiant; virucide; ophthalmological; keloid; skin disorder; Insulin-like Growth Factor I receptor; IGF-1; pitvriasis; IGF binding protein; IGFBP-2; IGFBP3; inflammation; psoriasis; pilaris; growth factor mediated cell proliferation; itothyosis; serborrhoea; ruba; keratosis; neoplasia; seleroderma; wart; skin cancer; sclerotic disease; hyperneovascular condition; hyperplasia; kidney disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Ameliorating the effects of a disorder, e.g. psoriasis, by administering UV (ultra-violet) treatment (optional) and an antisense nucleic acid that inhibits or reduces growth factor mediated cell proliferation and/or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               hyperneovascular condition; hyperplasia; kidney disease; neovascular condition of the retina; ss.
                                                                                                                                                                                                                                                    30-MAR-2001
                                                                                                                                                                                                                                                                                                                               AAF49085 standard; DNA; 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Example 8; Page 61; 201pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WPI; 2001-041421/05
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                                                                                                                                                                                                               IGF-I oligonucleotide #45
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   12;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Similarity
                                           condition of the retina;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            BP; 4 A;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Werther
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100.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            2 C;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0 U;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0;
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ART49619/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       receptor, IGF binding protein [IGFBP]-2 or IGFBP3], which is capable of inhibiting or reducing growth factor mediated cell proliferation, inflammation and/or other disorders. The present sequence is an oligonucleotide which can be used to design the antisense oligonucleotides of the present invention (see AAF45151 and AAF45153-F45161). The method is useful for amellorating the effects of psoriasis, ichthyosis, pityriasis, ruba, pilaris, serborrhoea, keloids, keratosis, neoplasias, scleroderma, warts, benign growths, cancers of the skin, a hyperneovascular condition such as a meovascular condition of the retina, brain or skin, growth factor-mediated malignancies, other sclerotic disease, kidney disease, hyperproliferation of the inside of blood vessels or any other hyperplasia
                                                                                                                                      neutral lipid transfer; plasma lipoprotein; atherosclerosis; atherectomy; reverse cholesterol transport; high density lipoprotein; therapy; CETP; familial hypercholesterolaemia; dyslipidaemia; hypoalphallpoproteinaemia; peripheral vascular disease; hyperbetalipoproteinaemia; RCT; inhibitor; angioplastic restenceis; low density lipoprotein; diabetes; HDL; human;
                                                                                                                                                                                                                            Hammerhead ribozyme; cholesterol ester transfer protein; mRNA cleavage;
                                                                                                                                                                                                                                                                                                28-FEB-1997
                                                                                                                                                                                                                                                                                                                                                                   AAT49619 standard; RNA; 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 15 BP; 4 A; 3 C; 4 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     skin disorders. The method comprises contacting the skin with antisense oligonucleotide, (for Insulin-like Growth Factor [IC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Example 8; Page 61; 201pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WPI; 2001-041421/05
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Wraight CJ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  21-JUN-2000; 2000WO-AU000693
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                                                                                      Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           inflammation.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           inhibits or reduces
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       1490 AGCCAGACTTCA 1501
                                                                                                                        S
                                                                                                                                                                                                                                                                                                                                                                                                                                                         14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        l Similarity
12; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                       AGCCAGACTTCA
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                                                                                                                                                                                                                                                                                              (first entry)
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                                                                                                                                                                                                                                                              ribozyme target sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             treatment
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         fects of a disorder, e.g. psoriasis, by admin
reatment (optional) and an antisense nucleic
growth factor mediated cell proliferation an
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              GA,
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                                                                                                                                                                                                                                                                                                                                                                   ВP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         0,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Score 12;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         to a method for ameliorating the effects
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Mismatches
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                                                                                                                                                                                                                                                              #477.
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18;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             e.g. psoriasis, by administering
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RESULT 16
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CC The HH ribozymes can also be used diagnostically to study genetic drift activity.
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Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             New ribozyme(s) for cleaving cholesterol ester transfer protein mRNA useful for preventing or treating initial development, progression or regression of vascular diseases, esp. familial hypercholesterolaemia.
                                                              Antisense therapy; antiproliferative; antiinflammatory; antipsoriatic; cytostatic; dermatological; cardiant; virucide; ophthalmological; keloid; skin disorder; Insulin-like Growth Factor I receptor; IGF-1; pityriasis; IGF binding protein; IGFB-2; IGFBB3; inflammation; psoriasis; pilaris; growth factor mediated cell proliferation; ichthyssis; serborrhoea; ruba; keratosis; neoplasia; scleroderma; wart; skin cancer; sclerotic disease; hyperneovascular condition; hyperplasia; kidney disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Claim 4;
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                                                                                                                                                                                                                                                     30-MAR-2001
                                                                                                                                                                                                                                                                                                                           AAF47839 standard; DNA; 15
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                                                   neovascular condition of the
                                                                                                                                                                                                                  IGFBP3
                                                                                                                                                                                                                                                                                                                                                                                                                                                      1488
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          13;
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                                                                                                                                                                                                                oligonucleotide #1259.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 BP; 4 A; 3 C; 4 G; 0 T; 4 U; 0 Other;
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                                                                                                                                                                                                                                                   (first entry)
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                                                                                                                                                                                                                                                                                                                           ВÞ
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Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         oligonucleotides of the present invention (see AAF45151 and AAF45153-F45161). The method is useful for ameliorating the effects of psoriasis, ichthyosis, pityriasis, ruba, pilaris, serborrhoea, keloids, keratosis, neoplasias, scleroderma, warts, benign growths, cancers of the skin, a hyperneovascular condition such as a neovascular condition of the retina, brain or skin, growth factor-mediated malignancies, other sclerotic disease, kidney disease, hyperproliferation of the inside of blood vessels or any other hyperplasia
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        skin disorders. The method comprises contacting the skin with an antisense oligonuclectide, (for Insulin-like Growth Factor [IGF]-1 receptor, IGF binding protein [IGFBP]-2 or IGFBP3), which is capable inhibiting or reducing growth factor mediated cell proliferation, inflammation and/or other disorders. The present sequence is an oligonuclectide which can be used to design the antisense
                                                                                                                                                                                                                                                                                   AAX57564;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence
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                                                     03-NOV-1998;
                                                                                                                                                                                                  Antisense; human; insulin-like growth factor-1 receptor; IGF-1R;
                                                                                                                                                                                                                           Antisense
                                                                                                                                                                                                                                                                                                              AAX57564
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Ameliorating the effects of a disorder, e.g. psoriasis, by administering UV (ultra-violet) treatment (optional) and an antisense nucleic acid that inhibits or reduces growth factor mediated cell proliferation and/or
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(INEX-) INEX PHARM CORP
                          04-NOV-1997;
                                                                                   14-MAY-1999
                                                                                                             WO9923259-A1
                                                                                                                                         Homo
                                                                                                                                                        Synthetic
                                                                                                                                                                                 expression;
                                                                                                                                                                                                                                                       16-JUL-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                The present invention relates to a method for ameliorating the effects
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        inflammation.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (MURD-)
                                                                                                                                                                                                                                                                                                                                                                                                                                                         Local
                                                                                                                                                                                                                                                                                                                                                                                                                1487
                                                                                                                                         sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                            13;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                        Similarity
                                                                                                                                                                                                                                                                                                              standard;
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                                                                                                                                                                                                                                                                                                                                                                                  AGCACCCAGACTTCA 15
                                                                                                                                                                                                                           oligo #3 to insulin-like growth factor I receptor
                                                                                                                                                                                 inhibition; induction; apoptosis; tumour; liposome; ss
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   BP; 5 A; 6 C; 2 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                            Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             2000WO-AU000693
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                                                      98WO-US023418
                                                                                                                                                                                                                                                                                                                DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                        39.3%;
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ARESULT 18
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                                                                     The present sequence represents a degenerate PCR primer for the M antig isolated from Histoplasma capsulatum (HC). HC polypeptides can be used for detecting antibodies for detecting a previous or current HC infectin a subject. They can also be injected into the skin of a subject to detect past exposure to HC by detecting swelling of the skin. The articles can be used for detecting current HC infection in a subject.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequences AAX57562-X57571 represent antisense oligonucleotides targeted to a region spanning 4-9 codons downstream of the AUG translation initiation codon of the human insulin-like growth factor-1 receptor (IGF-IR) gene. The antisense oligonucleotides inhibit the expression of IGF-
HC nucleic acids and polypeptides can also be used for the treatment of histoplasmosis as well as in vaccines for the prevention of histoplasmosis
                                                                                                                                                                                                                                                                                                                         New isolated Histoplasma capsulatum nucleic acids, used products for the diagnosis, prevention and treatment of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Histoplasma capsulatum; fungus; M antigen; vaccine; detection; histoplasmosis; diagnosis; infection; antimicrobial; antibody;
                                                                                                                                                                                                                                                                         Example 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                               Zancope-Oliveira RM,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         30-APR-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   04-NOV-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Synthetic
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WPI; 1999-313361/26
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            27-APR-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WO9955874-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Ajellomyces capsulatus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAZ35973;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (USSH ) US DEPT HEALTH & HUMAN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      primer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            which in turn induces apoptosis, especially in yonucleotides can be administered via a liposome
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                1488 GAAGCCAGACTTC 1500
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                                                                                                                                                                                                                                                                      Page 49; 75pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             BP;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                            Lott TJ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ΒP
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Score 11.4;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SERVICES
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histoplasmosis.
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Best Local S
Matches 10
                                                                  disease,
                                                                                     skin disorders. The method comprises contacting the skin with an antisense oligonucleotide, (for Insulin-like Growth Factor [IGF]-1 receptor, IGF binding protein [IGFBP]-2 or IGFB23), which is capable of inhibiting or reducing growth factor mediated cell proliferation, inflammation and/or other disorders. The present sequence is an oligonucleotide which can be used to design the antisense oligonucleotides of the present invention (see AAF45151 and AAF45153-F45161). The method is useful for ameliorating the effects of psoriasis, ichthyosis, pityriasis, ruba, pilaris, serborrhoea, keloids, keratosis, neoplasias, scleroderma, warts, benign growths, cancers of the skin, a hyperneovascular condition such as a neovascular condition of the retina, brain or skin, growth factor-mediated malignancies, other sclerotic disease, kidney disease, hyperproliferation of the inside of blood
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Antisense therapy; antiproliferative; antiinflammatory; antipsoriatic; cytostatic; dermatological, cardiant; virucide; ophthalmological; keloid; skin disorder; Insulin-like Growth Factor 1 receptor; IGF-1; pityriasis; IGF binding protein; IGFBP-2; IGFBP3; inflammation; psoriasis; pilaris; growth factor mediated cell proliferation; ichthyosis; serborrhee; ruba;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           The present invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Ameliorating the effects of a disorder, e.g. psoriasis, by administering UV (ultra-violet) treatment (optional) and an antisense nucleic acid that
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WPI; 2001-041421/05
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hyperneovascular condition; hyperplasia; kidney disease;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               reatment (optional) and an antisense nucleic acid growth factor mediated cell proliferation and/or
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           relates to a method for ameliorating the effects
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No. 23;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              of.
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Query Match

Sequence

15

BP;

2 A; 4 C; 6

G; 3 T; 0 U;

0 Other;

Score

11.4;

ВB 1;

Length 15

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RESULT 20
ANZ64816/C
ID ANZ648
XX ANZ648
XX ANZ648
XX ANZ648
XX ANZ648
XX Substr
XX Substr
XX Enzyma
KW Cirrho
KW autoim
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                                                                                                                       Query Match
Best Local Similarity
Matches 12; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Best Local Similarity Matches 12; Conserv
                                                                                                                                                                                                                                                                                                                                                                              The present sequence represents the preferred target sequence of an enzymatic nucleic acid, especially a hairpin ribozyme, which cleaves the Hepatitis C virus (HCV) RNA sequence at the base position given in the descriptor line. The HCV sequence was screened for optimal ribozyme target sites using a computer folding algorithm and regions of the mRNA which did not form secondary folding structures and contained potential ribozyme cleavage sites were identified. Ribozymes were synthesised to target these sites and their activities optimised by either varying the length of the binding arms or by modification to prevent degradation by nucleases. The ribozymes or by modification inhibit gene expression and/or viral replication, and are used to treat diseases associated with hepatocellular carcinoma. The ribozymes may be used in combination with inferting to the interference of the infertion of the interference of the infertion of the interference of the int
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         27-APR-1998;
18-SEP-1998;
25-FEB-1999;
                                                                                                                                                                                                                                                             Sequence
                                                                                                                                                                                                                                                                                                                           hepatocellular carcinoma. The ribozymes may be used in combin interferon to treat HCV infection, other infectious diseases, diseases, and cancer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Claim 2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Novel ribozymes for the treatment hepatitis C infection.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               04-NOV-1999.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Enzymatic nucleic acid; hammerhead ribozyme; virus replication; cleavag cirrhosis; liver failure; hepatocellular carcinoma; interferon; cancer; autoimmune disease; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Substrate
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           28-MAR-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AAZ64816 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              26-APR-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Hepatitis C virus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAZ64816;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (RIBO-)
                                                             1495
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1488 GAAGCCAGACTTC 1500
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14
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            RIBOZYME PHARM INC.
                                                          GACTTCAGCAGCCA 1508
                                                                                                                                                                                                                                                             14 BP;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Page 98; 123pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       for hairpin ribozyme which cleaves HCV at nt.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Mcswiggen
                                                                                                                              Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Conservative
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98US-0100842P.
99US-00257608.
99US-00274553.
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                                                                                                                                                                                                                                                          2 A; 5 C; 3 G; 0 T; 4 U;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       RNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     entry)
                                                                                                                                                          36.0%;
85.7%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         92.3%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Roberts
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                                                                                                                                                          Score 10.8;
Pred. No. 27;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Pred. No.
                                                                                                                              Mismatches
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                                                                                                                                                                                                                                                          0 Other;
                                                                                                                                                                                        DB
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           PA,
                                                                                                                                                                                        1;
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                                                                                                                                                                                        Length 14;
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                                                                                                                              Indels
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                                                                                                                       Gaps
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AAA99958/c
ID AAA999
XX AAA999
XC AAA999
XC Gemini
XX Facuou
PN W02000
PD 27-JUL
XX 26-JAN
PR 26-JAN
PR 26-JAN
PR 26-JAN
PR 27-JAN
PR 27
RESULT 22
ABX01653/c
ID ABX01653/c
XX ABX01
XX
AC ABX01
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C ABX01
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
Best Local
                                                                                                                                                          Enzymatic nucleic acid; RNA cleavage; Hepatitis C virus infection; HCV ribozyme; HCV expression; HCV replication; cirrhosis; virucide; liver failure; hepatocellular carcinoma; HCV infection; drug therapy; type I interferon; interferon alpha; interferon beta; cytostatic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                to geminivirus, involving introducing a geminivirus replication associated protein (Rep)-iteron antagonist into a plant. The antagonist is a nucleotide sequence of a geminivirus iteron capable of binding to a sequence of a geminivirus iteron capable or a defective Rep which has a conserved geminivirus iteron binding site. The present sequence is a geminivirus Rep repeat motif sequence. (Updated on 15-SEP-2003 to standardise OS field)
                                                                         type I interferon;
interferon gamma; c
substrate; hairpin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ABX01653;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 14 BP; 1 A; 3 C; 6 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Producing plants resistant to geminivirus, and inhibiting geminivirus replication in plants, by introducing replication associated protein iteron antagonists into the plant, plant cells or propagules.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Geminivirus
                                                                                                                                                                                                                                                                                                                                                                         Hepatitis C virus substrate #138 for HCV hairpin ribozyme
                                                                                                                                                                                                                                                                                                                                                                                                                                                               23-DEC-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ABX01653
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Example 1; Page 51; 172pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          26-JAN-1999;
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25-JAN-2001
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAA99958 standard; DNA; 14
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           27-JAN-2000; 2000WO-US001849.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              plant;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      The present invention relates to methods for producing plants resistant
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (SCRI)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1487 AGAAGCCAGACTTC 1500
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             2000-499224/44.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    l Similarity
12; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              88.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AGACGCCAGACTCC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Chatterji A;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Rep repeat motif sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (revised)
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                                                                                                                            consensus interferon; hepatotropic; antiinflammatory;
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85.7%;
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Pred. No. 27
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Hepatitis C virus

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RESULT 23
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ribozymes are useful for modulating the expression and/or replication of HCV. They can be used to treat cirrhosis, liver failure and/or hepatocellular carcinoma. The HCV ribozymes are also useful for treating a condition associated with HCV ribozymes are also useful for core other drug therapies, particularly type I interferon, especially interferon alpha, beta or gamma or consensus interferon. The present sequence represents a substrate for a HCV hairpin (HP) ribozyme. Note: Some of the sequence data for this patent did not form part of the printed specification. The complete sequence data for this patent was obtained in electronic format directly from the USPTO web site at
                                                                                              SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       The present invention relates to enzymatic nucleic acids which specifically cleave RNA derived from Hepatitis C virus (HCV). The enzymatic nucleic acid or ribozyme is in a hammerhead (HH) or hairpin (HP) motif where the binding arms comprise sequences complementary to of the substrate sequences defined in the specification. The HCV of the substrate sequences defined in the specification.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          New ribozymes targeting RNA derived from hepatitis C virus inhibit viral replication and are useful to treat hepatitis C virus infections and cirrhosis, liver failure or hepatocellular carcinoma.
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(PAVC/)
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               18-OCT-2001
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                                                                                                                                                                                                                  ABF18234;
                                                                                                                                                                                                                                               ABF18234 standard; DNA;
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                                                                     Homo sapiens
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PAVCO P A.
MACEJACK D.
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85.7%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               07-APR-2000; 2000DE-01019173
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SNP; single nucleotide polymorphism; human;
peptide nucleic acid; cytosine methylation;
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Set of oligonucleotides, useful for diagnosis and cell typing, designed to detect single-nucleotide polymorphisms and cytosin

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RESULT 25
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                                                                                                                                                                                                                                                                          19-JUN-1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SAGE tag; serial analysis of gene expression; antigen-presenting ceAPC; monocyte-derived dendritic cell; differential gene expression; immunostimulatory cofactor; costimulatory factor; CTL;
                                                                                                                                                                         19-JUN-1998;
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98US-0090040P.
98US-0090041P.
98US-0090043P.
98US-0090044P.
98US-0090044P.
98US-0090047P.
98US-0090047P.
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98US-0090035P.
98US-0090036P.
98US-0090039P.
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98US-0089844P.
98US-0089853P.
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98US-0089991P
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                                                                                                                                                                                                                                                                                                                                                  -0089997P
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Score 10.4;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Pred.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SEQ ID NO:1327.
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RESULT 26
AAZ83653/c
ID AAZ836
XX
AC AAZ836
AC AAZ836
XX
DT 07-APR
XX
DE Metast

AAZ83653 standard; DNA; 10

07-APR-2000 AAZ83653;

(first entry)

Metastatic breast tumour cell upregulated transcript tag #2887.

뮍 S

 $\vdash$ 

AGAAGCCAGA 10

1487 AGAAGCCAGA 1496

Query Match Best Local S Matches 10

Similarity

ilarity 100.0%; I Conservative 0;

Mismatches

0,

Gaps

0

33.3%; Score 10; 100.0%; Pred. No.

DB 1; Length 10; 28;

10;

Sequence

10

B₽;

v A;

2 C; 3

G; 0 T; 0

u; o

Other;

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CC complex) and subsequent recognition by T-cell receptors is alone insufficient to activate a robust cytotoxic immune response that can lyse the tumour cells, immunostimulatory cofactors also being required for efficient activation of cytotoxic T-lymphocytes (CTLs). Nucleic acid for efficient activation of cytotoxic T-lymphocytes (CTLs). Nucleic acid sequences identified using the SAGE tags have several potential uses. They may be used in vaccines to induce an immune response, particularly against a tumour antigen; to modulate the genotype of an APC; to screen for agents that modulate expression of differentially expressed genes in APC; and as hybridisation probes/amplification primers for the charges of these genes. Detection of the dendritic cell differentially expressed genes, or of their encoded proteins, can be used to identify cells as belonging to the monocyte lineage. Cells containing these genes can be used in active immunotherapy (or to stimulate production of a population of antigen-specific effector cells) and vectors containing these genes can be used in gene therapy. Co-administration of tumour antigens and apresentation to endogenous APCs and upregulates the APCs for the presentation of co-stimulatory signals, migration to T cell-rich sites, secretion of T cell growth factors and secretion of chemokines for cervitment of immune effector cells
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      19-JUN-1998;
19-JUN-1998;
19-JUN-1998;
19-JUN-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            with monocytes. Some of the transcripts correspond to known genes or ESTE (expressed sequence tags) which were previously unknown to be preferentially or differentially expressed in dendritic cells, while other transcripts correspond to novel genes. Antigen-presenting cell (APC)-associated costimulatory factors play an important role in the activation of the cytotoxic immune response, particularly against tumour cells. Tumour antigen presentation via the MHC (major histocompatibility
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequences AAZ77573-Z79709 represent SAGE (serial analysis of gene expression) tags used to identify mRNA transcripts encoding immunostimulatory cofactor proteins which are preferentially or differentially expressed in monocyte-derived dendritic cells compared
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Claim 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         cells, useful in gene vaccines against cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Isolated polynucleotides differentially expressed in antigen-presenting
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Page 103; 130pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Shankara
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98US-0090077P.
98US-0090078P.
98US-0090079P.
98US-0090080P.
98US-0111715P.
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ESTS

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AAF39527
ID AAF3
XX
AC AAF3
XX
DT 23-N
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                                                                                                         RESULT 27
                                                                                                                                                                                                                                                                                                                                                 CC AAZ80767 to AAZ83941 represent tags corresponding to distinct transcripts CC that are preferentially transcribed in the metastatic breast tumour cells). AAZ83942 tissue (i.e. are upregulated in metastatic breast tumour cells) that are preferentially transcribed in the metastatic breast tumour cells). AAZ83942 CC to AAZ86677 represent tags corresponding to distinct transcripts that are preferentially transcribed in the primary or non-metastatic breast tumour cells). These creations is compounded to the primary or non-metastatic breast tumour cells). These cc transcripts can be used for diagnosis, prognosis, monitoring and creatment of breast cancer, particularly where metastatic. Diagnosis is by standard immunoassays or hybridisation/amplification reactions. CC compounds that modulate expression of the transcripts are potentially cuseful for treatment of (metastatic) breast cancer, while promoters from ct the transcripts are used to direct expression, in selected cell types, of e.g. therapeutic genes (also ribozymes or antisense sequences), of particularly an antigen-encoding sequence for use in gene or cell-based corrections; for diagnosing breast cancer and for ratsing specific cantibodies (Ab). Ab are used to detect the polypeptides or as therapeutic and isolate populations of educated, antigen-specific immune effecter cells, e.g. cytotoxic T lymphocytes, and these used for adoptive
                                                                                                                                                                                                                                                 Query Match
Best Local :
                                                                                                                                                                                                                                 Best Local Similarity Matches 10; Conserv
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19-JUN-1998;
19-JUN-1998;
19-JUN-1998;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human; metastatic breast tumour tissue; breast cancer; tag; primer; non-metastatic breast tumour tissue; gene therapy; anticancer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Isolated polynucleotides differentially expressed between metastatic non-metastatic breast cancer cells, useful for diagnosis, prevention
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                                    AAF39527;
                                                                                                                                                                                                                                                                                                     Sequence 10 BP; 2 A; 2 C; 3 G; 3 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     treatment of cancer.
 23-MAR-2001
                                                                       AAF39527 standard; DNA; 10
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ROBERTS B L.
SHANKARA S.
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 (first entry)
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98US-0089997P.
98US-0090039P.
98US-0090040P.
98US-0090041P.
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                                                                                                                                                                                                 33.3%; occ. No. -
100.0%; Pred. No. -
'Ave 0; Mismatches
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Yeast NORF gene SAGE tag oligonucleotide SEQ ID NO:6266
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Yeast; Saccharomyces cerevisiae; characterisation; cell cycle; NORF; nor previously assigned open reading frame; nonannotated ORF; SAGE; serial analysis of gene expression; antifungal; tag; identification; linker; PCR primer; ds.

Saccharomyces cerevisiae.

WO200077214-A2

14-JUN-2000; 2000WO-US016223

16-JUN-1999; 99US-00335032

(UYJO ) UNIV JOHNS HOPKINS

Velculescu V, Vogelstein B, Kinzler K;

WPI; 2001-061874/07.

Yeast gene coding sequences comprising NORF genes with serial analysis gene expression (SAGE) tags, useful for studying, monitoring and affecting phases of the cell cycle. O.f

Example; Page 223; 419pp; English.

Coding sequence of a yeast gene selected from a group of 745 NORF (not coding sequence of a yeast gene selected from a group of 745 NORF (not comprising a ssigned open reading frame, or nonannotated ORF) genes comprising a SAGE (serial analysis of gene expression) tag. Also codescribed are: (1) a method (M1) of using NORF genes to affect the cell cycle comprising administering a NORF gene whose expression varies by at cell east 10% between any two phases of the cell cycle selected from log cantifungal drugs comprising: (a) a method (M2) for screening candidate cell; and (b) monitoring expression of a NORF gene whose expression of the yeast gene is a candidate antifungal drug; (3) a method (M3) for cell; and (b) monitoring expression of a NORF gene whose expression of the yeast gene is a candidate antifungal drug; (3) a method (M3) for comprising contacting human DNA with a probe which comprises at least 10 comprising contacting human but whose expression varies as in M1; contiguous nucleotides of a NORF gene whose expression varies as in M1; cell cycle progression in the yeast cell of at least 10 comprising contacting a yeast cell with a candidate drug as a member of a class of drugs having a characteristic effect on gene expression in a yeast cell with a candidate drug and member of a cycle and (a) a method (M4) for identifying a candidate drug as a member of a cycle and spression in the yeast cell of at least 1 NORF gene whose expression in the yeast cell of the cell cycle. The method whose candidate drug and my be used to identify candidate drugs. AAF33268 to AAF33267 represent 1inkers and PCR primers used in the SAGE cycle and for identification of the present invention.

CC AAF33262 to AAF33267 represent linkers and PCR primers used in the SAGE method, in the exemplification of the present invention.

Sequence 10 BP; 5 A; 2 C; 3 G; 0 T; 0 U; 0 Other;

Ś 밁 Query Match Best Local Similarity Matches 10; Conserv 1487  $\vdash$ AGAAGCCAGA 1496 Conservative 100.0%; 33.3%; 0; Score 10; Pred. No. Mismatches DB 1; 28; 0 Length 10 0

Gaps

0

RESULT 28
AAF34162
ID AAF3
XX

AGAAGCCAGA

AAF34162 standard; DNA; 10 BP

Yeast NORF gene SAGE tag oligonucleotide SEQ ID NO:901

23-MAR-2001 (first entry)

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                                      δ
                                                                                                                                                                     CC Coding sequence of a yeast gene selected from a group of 745 NORF (not CC previously assigned open reading frame; or nonamnotated ORF) genes CC comprising a SAGE (serial analysis of gene expression) tag. Also CC comprising a SAGE (serial analysis of gene expression) tag. Also CC described are: (1) a method (M1) of using NORF genes to affect the cell CC cycle comprising administering a NORF gene whose expression varies by at CC least 10% between any two phases of the cell cycle selected from log CC phase, S phase and G2/M; (2) a method (M2) for screening candidate CC cell; and (b) monitoring expression of a NORF gene whose expression of CC varies as in M1, where a test substance which modifies the expression of CC identifying human genes which are involved in cell cycle progression CC comprising contacting human DNA with a probe which comprises at least 10 contacting a candidate drug; (3) a method (M3) for CC contiguous nucleotides of a NORF gene whose expression varies as in M1; and (4) a method (M4) for identifying a candidate drug as a member of a CC class of drugs having a characteristic effect on gene expression in a CC class of drugs having a characteristic effect on gene expression in a CC class of drugs having a characteristic effect on gene expression in a CC contiguous nucleotides of a NORF gene whose expression in a contacting expression in the yeast cell with a candidate drug and CC monitoring expression in the yeast cell of at least 1 NORF gene whose expression is affected by the class of drugs. The NORF genes may be used to study, monitor and affect phases of the cell cycle, the differentially captions of the coll cycle and for identification of antifungal drugs which affect the cell cycle. The CC methods may be used in the exemplification of the present invention. CC AAF32267 represent linkers and PCR primers used in the SAGE contaction of the present invention.
                                                                                     Matches
                                                                                                                        Query Match
                                                                                                                                                                   Sequence 10 BP; 5 A; 2 C; 3 G; 0 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   affecting phases of the cell cycle.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Yeast gene coding sequences comprising NORF genes with serial analysis gene expression (SAGE) tags, useful for studying, monitoring and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                14-JUN-2000; 2000WO-US016223
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                                        1486 AAGAAGCCAG 1495
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ш
                                                                                                         Similarity
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                                                                                                                                                                                                           the exemplification of the present invention
                                                                                  Conservative
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                                                                                                    100.0%;
                                                                                                                           33.3%;
                                                                                  0
                                                                                                       Score 10;
Pred. No.
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                                                                                  Mismatches
                                                                                                       DB 1;
. 28;
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                                                                                                                        Length 10
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                                                                              Gaps
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RESULT 29

밁 Ś

Matches

10;

1496 ACTTCAGCAG 1505

Local

Similarity

larity 100.0%; Conservative

Score 10; Pred. No.

DB 1; Length 10, 28;

Mismatches

0

Indels

0

Gaps

0

Sequence 10

BP; 3

A; 3 C; 2 33.3%;

G; 2 T; 0 U; 0 Other;

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cc phase, S phase and G2/M; (2) a method (M2) for screening candidate CC phase, S phase and G2/M; (2) a method (M2) for screening candidate artifungal drugs comprising: (a) contacting a test substance with a yeast cell; and (b) monitoring expression of a NORF gene whose expression of CC they east gene is a candidate antifungal drug; (3) a method (M3) for CC identifying human genes which are involved in cell cycle progression of CC contiguous nucleotides of a NORF gene whose expression varies as in M1; CC and (4) a method (M4) for identifying a candidate drug as a member of a CC class of drugs having a characteristic effect on gene expression in a CC yeast cell comprising contacting a yeast cell with a candidate drug and CC expression is affected by the class of drugs. The NORF gene whose cexpression is affected by the class of drugs. The NORF gene may be used CC expressed genes may be used as markers of phases of the cell cycle. The methods may be used as markers of phases of the cell cycle. The CC methods may be used as markers of phases of the cell cycle. The CC cycle and for identification of antifungal drugs. AAF33268 to AAF33267 represent linkers and PCR primers used in the same scene invention.

CC AAF33262 to AAF33267 represent linkers and PCR primers used in the SAGE
                                                                                                                                                                                                                                                                                                                                                                                                                                                      The present invention describes an isolated DNA molecule comprising a coding sequence of a yeast gene selected from a group of 745 NORF (not previously assigned open reading frame, or nonannotated ORF) genes comprising a SAGE (serial analysis of gene expression) tag. Also described are: (1) a method (M1) of using NORF genes to affect the cell cycle comprising administering a NORF gene whose expression varies by at least 10% between any two phases of the cell cycle selected from log phase, S phase and GZ/M; (2) a method (M2) for screening candidate
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Yeast gene coding sequences comprising NORF genes with serial analysis of gene expression (SAGE) tags, useful for studying, monitoring and affecting phases of the cell cycle.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 21-DEC-2000
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Yeast; Saccharomyces cerevisiae; characterisation; cell cycle; NORF; nor previously assigned open reading frame; nonannotated ORF; SAGE; serial analysis of gene expression; antifungal; tag; identification;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Yeast NORF gene SAGE tag oligonucleotide SEQ
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAF39463 standard; DNA; 10 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Example; Page 221; 419pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           16-JUN-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    14-JUN-2000; 2000WO-US016223
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (UYJO ) UNIV JOHNS HOPKINS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             PCR primer; ds.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Vogelstein B,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             99US-00335032
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Kinzler K;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            IJ
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RESULT 30
AAF41899
                                                                                       Best Local Similarity
                                             Query Match
        Matches
                                                                                                                                                                                                                                                                                                                                                                                                                previously assigned open reading frame; or nonannotated ORF) genes comprising a SAGE (serial analysis of gene expression) tag. Also described are: (1) a method (M1) of using NORF genes to affect the cell cycle comprising administering a NORF gene whose expression varies by at least 10% between any two phases of the cell cycle selected from log phase, S phase and G2/M; (2) a method (M2) for screening candidate antifungal drugs comprising: (a) contacting a test substance with a yeast cell; and (b) monitoring expression of a NORF gene whose expression of aries as in M1, where a test substance which modifies the expression of the yeast gene is a candidate antifungal drug; (3) a method (M3) for identifying human genes which are involved in cell cycle progression.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     nor previously assigned open reading frame; nonannotated ORF; SAGE; serial analysis of gene expression; antifungal; tag; identification; linker; PCR primer; ds.
                                                                                                                                                                                                       comprising contacting human DNA with a probe which comprises at least 10 contiguous nuclectides of a NORF gene whose expression varies as in M1; and (4) a method (M4) for identifying a candidate drug as a member of a class of drugs having a characteristic effect on gene expression in a yeast cell comprising contacting a yeast cell with a candidate drug and monitoring expression in the yeast cell of at least 1 NORF gene whose expression is affected by the class of drugs. The NORF genes may be used to study, monitor and affect phases of the cell cycle, the differentially expressed genes may be used as markers of phases of the cell cycle. The methods may be used to identify candidate drugs which affect the cell methods may be used to identify candidate drugs which affect the cell
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAF41899
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAF41899 standard;
                                                                                                                            cycle and for identification of antifungal drugs. AAF33268 to AAF44064 represent SAGE tags used in the exemplification of the present invention. AAF33262 to AAF33267 represent linkers and PCR primers used in the SAGE method, in the exemplification of the present invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Yeast gene coding sequences comprising NORF genes with serial analysis of gene expression (SAGE) tags, useful for studying, monitoring and affecting phases of the cell cycle.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 2001-061874/07.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Velculescu V,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WO200077214-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Saccharomyces cerevisiae
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Yeast; Saccharomyces cerevisiae; characterisation; cell cycle; NORF;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Yeast NORF gene SAGE tag oligonucleotide SEQ ID NO:8638
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         23-MAR-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    The present invention describes an isolated DNA molecule comprising a coding sequence of a yeast gene selected from a group of 745~\mathrm{NORF} (not
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Example; Page 308; 419pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        14-JUN-2000; 2000WO-US016223
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                                                                                     Sequence 10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (UYJO ) UNIV JOHNS HOPKINS
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        10;
                                                                                       BP;
      Conservative
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100.0%; --
                                                                                     2 C; 3 G; 0 T; 0 U; 0 Other;
                       33.3%; Score 10;
100.0%; Pred. No.
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        Mismatches
                           28;
                                               DB 1; Length 10;
        0
        Indels
        0;
        Gaps
        0
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RESULT 31
ANV32260/c
ID ANV322
XX ANV322
XX ANV322
XX ANV322
XX ANV322
XX RT-PCR
XW differ
XX Synthe
XX C-APR
X
RESULT 32
AAX89228
ID AAX89
XX
AC AAX89
XX
DT 15-SE
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                                                                                                                                                                                                                                                                               B
                                                                                                                                                                                                                                                                                                                                                                                                    Best Local Similarity Matches 10; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      expression by random primed reverse transcription PCR (RT-PCR). The primer sequences are generated by stimulating PCR reactions on non-redundant mammalian nucleotide sequence databank entries containing at least 1,000 bp of coding region. The primers selected, such as the present one, had to meet various criteria such as having an efficiency index between 2-10, having a selectivity index higher than 1, being 12 bp long i.e. 8 C or G and 4 T or A, and each primer differed from the others in at least 5 of the 8 bases at the 3'-end. The invention claims the selected primers make it possible to use internally primed, PCR-based RNA fingerprinting for simple, exhaustive and systematic analysis of differential gene expression as an advantageous alternative to differential display. The method can also be useful for isolating new coding sequences and to compare known and new genes
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               differential gene expression; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     RT-PCR; primer; amplification; reverse transcription; RNA fingerprinting;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Random primed reverse transcription PCR primer 108
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAV32260
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAV32260 standard; DNA; 12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Differential screening of gene expression by reverse transcription polymerase chain reaction - uses random priming with primers selected for high efficiency and selectivity by computer screening of database(s).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              02-APR-1998
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 12 BP; 0 A; 3 C; 4 G; 4 T; 0 U; 1 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    The invention provides a method for the differential screening of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Claim 9; Page 24; 37pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 1998-230725/20..
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Consalez G,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      27-SEP-1996;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  26-SEP-1997;
                                                                                                                           AAX89228
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (SANR-) FOND
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                                                                                                                                                                                                                                                                                                                            1479 CACGACCAAG 1488
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                                                                                                                              standard; DNA; 13
                                                                                                                                                                                                                                                                               CACGACCAAG
                                                                                                                                                                                                                                                                                                                                                                                                        Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Fesce R;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                    100.0%;
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                                                                                                                                                                                                                                                                                                                                                                                                        0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                    Score 10;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                           0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Length 12
                                                                                                                                                                                                                                                                                                                                                                                                           Indels
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                                                                                                                                                                                                                                                                                                                                                                                                           Gaps
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15-SEP-1999 AAX89228;

(first entry)

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                                                                                                                                                                                                                                                                                                                                                                                                The invention provides an antibody specific for epitopes (AAY27164-175) located on the extracellular domain of STZL protein. The antibody is used: (a) in a binding assay to identify and/or distinguish T-helper 1 (Th1) or Th2 and evaluate the abundance of Th2 in a sample of body fluid or tissue; (b) to lyse Th2 cells in preference to Th1 cells or otherwise inhibit Th2 cell function; and (c) as a medicament for use in therapy as a mediator of an allergic reaction, e.g. asthma; diseases mediated by intracellular pathogens, especially arthritis. The antibody is targeted to an epitope (STZL) which is a member of the immunoglobulin superfamily and is found on Th2 cells and not Th1 cells
                                                                                            Synthetic
                                                                                                                    Camellia sinensis L. (O.) Kuntze;
                                                                                                                                            Camellia
                                                                                                                                                                                                                  ADC64953 standard; DNA; 13
                                                                                                                                                                                                                                                                                                                                                                           Sequence 13 BP; 4 A; 4 C; 3 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            24-DEC-1998;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Synthetic
          30-MAR-2001; 2001CN-00112459
                                  06-NOV-2002
                                                          CN1377966-A
                                                                                 Camellia sinensis.
                                                                                                                                                                    18-DEC-2003
                                                                                                                                                                                           ADC64953;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Disclosure; Page 12; 44pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Novel
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WPI; 1999-430265/36
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ST2L; extracellular domain; mouse; epitope; antibody; T-helper cell; Th1 Th2; medicament; therapy; allergic reaction; asthma; Leishmania; intracellular pathogen; inflammatory disease; arthritis; immunoglobulin;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    PCR primer H-AP34.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ovel antibody to ST2L,
helper 1 cells.
                                                                                                                                                                                                                                                                                                      1497
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                                                                                                                                                                                                                                                                                           CTTCAGCAGC 1506
                                                                                                                                            sinensis L.
                                                                                                                                                                                                                                                                              CTTCAGCAGC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Xu D;
                                                                                                                                                                                                                                                                                                                              Conservative
                                                                                                                                                                  (first
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                                                                                                                                                                   entry)
                                                                                                                                                                                                                                                                              13
                                                                                                                                            (O.) Kuntze
                                                                                                                                                                                                                                                                                                                                         100.0%;
                                                                                                                                                                                                                                                                                                                                                    33.3%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    an epitope found
                                                                                                                                                                                                                    ВP
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                                                                                                                                                                                                                                                                                                                                         Score 10; DB : Pred. No. 36;
                                                                                                                                                                                                                                                                                                                              Mismatches
                                                                                                                    tea tree;
                                                                                                                                        related PCR primer AP34.
                                                                                                                                                                                                                                                                                                                                                   DB 1;
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                                                                                                                  PCR primer; ss
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                                                                                                                                                                                                                                                                                                                            0,
                                                                                                                                                                                                                                                                                                                                                   Length 13;
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RESULT 34
ABH65692/c
ID ABH656
XX ABH656
XX ABH656
XX SNP; R
KW Peptid
KW Centra
XX Homo R
XX Claim
XX This
CC Acid
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CC Cantr
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Best Local Similarity
Matches 10; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Cloning of a new gene sequence expressed and inhibited during winter dormancy of a tea tree top plumelet, comprises identification, cloning and analysis of a new primer in the gene sequence.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Oligonucleotide
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   The present invention describes the cloning of a new gene sequence expressed and inhibited during hibernation of the top plumelet of Camellia sinensis L.(O.) Kuntze tea tree. Also described is the
                                                                                                                                                                                                                                                                                  07-APR-2000;
                                                                                                                                                                                                                                                                                                                                                                         WO200177384-A2
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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory,

central nervous system, cardiovascular and

disorders.

Set of oligonucleotides, useful for diagnosis and cell designed to detect single-nucleotide polymorphisms and

typing, i

methylation

WPI; 2001-657177/75.

(EPIG-)

EPIGENOMICS

AG

Piepenbrock C,

Berlin

Claim 1; SEQ ID NO 265669; 29pp + Sequence Listing; German.

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RESULT 35
ABH65694/c
ID ABH656
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                                                                                                                                                     acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cyrosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for disgnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation, ABC00010-ABC9989, ABF00010-ABF9989, ABH00010-ABH99989 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at ftp.wipo.int/pub/published_pct_sequences
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    oligomers are also used for detecting cell type differentiation. ABC000 -ABC99989, ABF00010-ABF99989, ABH00010-ABF99989, ABH00010-ABF99989 and ABI00010-ABF99989 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, by the obtained in electronic format from WIPO at
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Set of oligonucleotides, useful for diagnosis and cell typing, designed to detect single-nucleotide polymorphisms and cytosine methylation status.
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                                                                                                               Sequence 13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   This invention describes novel oligonucleotide primers or peptide nucleic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Claim 1; SEQ ID NO 265671; 29pp + Sequence Listing; German.
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                                                                                                               BP;
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                                                                                                               0 Other;
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                                                                                                                                                                                                                                                                                                                     Sequence 13
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   07-APR-2000; 2000DE-01019173
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                                                                                                                                                                                                                                                                                                                     5 A;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AG.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       entry
                                                                                                                                                                                                    32.7%;
                                                                                                                                                                                                                                                                                                                     5
C;
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                                                                                                                                                                                                                                                                                                                         0
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                                                                                                                                                                                                                                                                                                                     G; 3 T; 0 U;
                                                                                                                                                              0;
                                                                                                                                                                                                    Score 9.8; I
Pred. No. 39;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ζ.
                                                                                                                                                                   Mismatches
                                                                                                                                                                                                                                                                                                                         0 Other;
                                                                                                                                                                                                                                          DB 1;
                                                                                                                                                              2
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                                                                                                                                                                                                                                      Length 13;
                                                                                                                                                                   Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            cancer; CNS;
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RESULT 38
ABQ83680/c
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Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                   Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010 -ABC99389, ABF00010-ABF99389, ABH00010-ABH99389 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this pattent did not form part of the printed specification, but was obtained in electronic format from WIPO at
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic
                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 13 BP; 6 A; 5 C; 0 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Set of oligonucleotides, useful for diagnosis and cell typing, designed to detect single-nucleotide polymorphisms and cytosine methylation status.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Olek A,
            19-MAR-2002; 2002WO-US008546
                                                                                                                             synthesis;
                                                                                                                                              Molecular function;
                                                                                                                                                                         DNA-templated
                                                                                                                                                                                                       27-JAN-2003
                                                                                                                                                                                                                                  ABQ83680
                                                                                                                                                                                                                                                                ABQ83680 standard; DNA; 13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Claim 1; SEQ ID NO 265672; 29pp + Sequence Listing; German.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WPI; 2001-657177/75
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 07-APR-2000; 2000DE-01019173
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             06-APR-2001; 2001WO-IB000713.
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                                         26-SEP-2002.
                                                                   WO200274929-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (EPIG-)
                                                                                                                                                                                                                                                                                                                                                                   1489 AAGCCAGACTTCA 1501
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                                                                                                                                                                                                                                                                                                                                        AACCCAAACTTCA 13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Piepenbrock C,
                                                                                                                             unction; diversification; selection; amplification; evolve; library; chemical compound; hybridisation; ss.
                                                                                                                                                                                                                                                                                                                                                                                                 Conservative
                                                                                                                                                                                                     (first entry)
                                                                                                                                                                      synthesis related oligonucleotide
                                                                                                                                                                                                                                                                                                                                                                                                                 84.6%;
                                                                                                                                                                                                                                                                                                                                                                                                                             32.7%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        265672 for detecting SNP TSC0064388.
                                                                                                                                                                                                                                                                ВP
                                                                                                                                                                                                                                                                                                                                                                                                 0
                                                                                                                                                                                                                                                                                                                                                                                                              Score 9.8; D
Pred. No. 39;
                                                                                                                                                                                                                                                                                                                                                                                                 Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                             DB 1; Length 13;
                                                                                                                                                                                                                                                                                                                                                                                                2
                                                                                                                                                                                                                                                                                                                                                                                                 Indels
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                                                                                                                                                                                                                                                                                                                                                                                                0
                                                                                                                                                                                                                                                                                                                                                                                                 Gaps
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contacting one or more transfer units having an anti-codon and reactive unit with the one or more templates under conditions to allow for the reactive units. Also described: (1) a method (M2) of evolving a library of compounds; (2) a kit comprising one or more non-natural polymers; and (4) a library (1) comprising one or more chemical compounds where each of the chemical compounds is bonded to an amplifiable template whose nucleotide sequence is informative of the structure of the chemical compounds where the synthesising one or more chemical compounds which contain an anti-codon comprising a nucleotide sequence which hybridises with one or more chemical compounds which contain an anti-codon comprising a nucleotide sequence which hybridises with one or more chemical compounds which contain an anti-codon comprising a nucleotide sequence which hybridises with one or more nucleic acid templates. M1 is useful for synthesising a library of chemical compounds that are not, or do not resemble nucleic acids or nucleic acid analogues. The present sequence represents an oligonucleotide which is
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Synthesizing chemical compounds by hybridizing one or more templates which have associated reactive unit, with one or more transfer units having anti-codon and reactive unit, and performing reaction of reactive
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        19-MAR-2001; 2001US-0277081P.
19-MAR-2001; 2001US-0277094P.
20-JUL-2001; 2001US-0306691P.
19-MAR-2002; 2002US-00101030.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Example 5; Fig 32; 146pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WPI; 2002-740858/80.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Liu
                                                                                                                                                                                                                                                                                                                                                                                                                                                             more chemical compounds. M1 involves providing one or more temp which optionally have a reactive unit associated with them; and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      units.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (HARD ) HARVARD COLLEGE.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           present invention describes a method (M1) for synthesising one or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Gartner ZJ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   or more templates,
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                    S
                                            Query Match

Best Local Similarity

Matches 11; Conserv
                                                                                        Sequence 13 BP; 3 A; 4 C; 3 G; 3 T; 0 U; 0 Other;
                      1491 GCCAGACTTCAGC 1503
                                          11;
13
 GGCAGATTTCAGC 1
                                             Conservative
                                                      32.7%;
84.6%;
                                             0
                                                      Score 9.8; I
Pred. No. 39;
                                                                    ВB
                                                                  1; Length 13
                                             Indels
                                             ٥,
                                            Gaps
                                             0
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RESULT 39
ABV70346/c
ID ABV703
XX ABV703
AC ABV703
XX J-OCT
XX 21-OCT
XX Human;
XX Human;
XX immuno
XX Imm
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       21-OCT-2002
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   psoriasis; dermatitis; skin cancer; EST; expressed sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Human; skin; dermatological; vulnerary; antipsoriatic; antiseborrhaeic; immunosuppressive; antiinflammatory; cytostatic; SAGE; neurodermatitis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ABV70346;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                mmunosuppressive;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 sapiens.
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20-DEC-2001; 2001WO-EP015179

11-JUL-2002.

03-JAN-2001; 2001DE-01000127

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RESULT 40
ABV69357/C
ID ABV69
XX ABV69
XX ABV69
XX Human
XX Human
XX Human
XX Immun
XX Immun
XX OS Homo
PN WO20(
XX WO20(
XX WO21)
PF 20-Di
XX O3-Ji
XX WPI;
XX WPI;
XX WPI;
XX WPI;
XX Disc
PS Disc
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Best Local
                                                                                                                                                                                                                                                                                                                                  Human; skin; dermatological; vulnerary; antipsoriatic; antiseborrhaeic; immunosuppressive; antiinflammatory; cytostatic; SAGE; neurodermatitis; psoriasis; dermatitis; skin cancer; EST; expressed sequence tag; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       The invention relates to in vitro identification (M1) of genes expressed in the skin of humans or animals by subjecting a mixture of genetically encoded factors from skin, to serial analysis of gene expression (SACE) so as to identify skin-expressed genes and quantify their expression. (M1) is useful for identifying genes involved in skin homeostasis; to determine skin homeostasis and to test agent (A) that maintains or promotes skin homeostasis or that can be used for treating skin.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         e.g. skin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 11 BP; 1 A; 3 C; 3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             disorders, specifically neurodermatitis; sunburn; psoriasis; scleroderma; ichthyosis; atopic dermatitis; acne; seborthea; lupus erythematosus; rosacea; melanoma; basal cell carcinoma; and carcinoma or sarcoma of the skin. The present sequence is that of a human expressed sequence tag
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Claim 24;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        In vitro identification of skin-expressed genes, useful for determining homeostasis and identifying cosmetic or pharmaceutical agents against
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WPI; 2002-590638/63
                             e.g. skin cancer
                                           In vitro identification of skin-expressed genes, useful for determining homeostasis and identifying cosmetic or pharmaceutical agents against
                                                                                          WPI; 2002-590638/63
                                                                                                                                                                                03-JAN-2001; 2001DE-01000127
                                                                                                                                                                                                              20-DEC-2001; 2001WO-EP015179
                                                                                                                                                                                                                                            11-JUL-2002.
                                                                                                                                                                                                                                                                          WO200253774-A2
                                                                                                                                                                                                                                                                                                       Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                              Human skin EST 7143
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ABV69357 standard; cDNA; 11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Petersohn D,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (HENK ) HENKEL
                                                                                                                                                    (HENK ) HENKEL KGAA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             1495 GACTTCAGCAG 1505
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         10;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               of the invention
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Page 259; 1345pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                           (first entry)
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                                                                                                                        Conradt
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                                                                                                                        Hofmann
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                German.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         0,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Pred.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Score 9.4;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          No.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       11;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         0
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in the skin of humans or animals by subjecting a mixture of genetically encoded factors from skin, to serial analysis of gene expression (SAGE) so as to identify skin-expressed genes and quantify their expression. (MI) is useful for identifying genes involved in skin homeostasis; to determine skin homeostasis and to test agent (A) that maintains or promotes skin homeostasis or that can be used for treating skin disorders, specifically neurodermatitis; sunburn; psoriasis; scleroderma; ichthyosis, atopic dermatitis; acne; seborrhea; lupus exythematosus; rosacea; melanoma; basal cell carcinoma; and carcinoma or sarcoma of the skin. The present sequence is that of a human expressed sequence tag (EST) of the invention
Sequence 11 BP; 0 A; 4 C; 2
     G; 5 T; 0 U; 0 Other;
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Query Match Best Local S Matches 10 1485 CAAGAAGCCAG 1495 10; Similarity Conservative 31.3%; 90.9%; Score 9.4; Pred. No. 4 Mismatches

DB 1; 1,

Length 11; Indels

0

Gaps

0

밁 8

11 CAAGAAGGCAG 1

ARESULT 41
ABV6295/C
ID ABV629
XX ABV629
XX ABV629
XX DT 21-OCT
XX Human;
KW Human;
KW Human;
KW Homo 8
XX WO2002
XX IN-JUL
XX O3-JAN
XX PA (HENK
XX D1-JAN
XX D1-JAN
XX D1-JAN
XX O3-JAN
XX D1-JAN
XX O3-JAN
XX PA (HENK
XX PA (HENK
XX D1-JAN
XX O3-JAN
XX O3-JAN
XX O3-JAN
XX PA (HENK
XX D1-JAN
XX O3-JAN
XX O3-Human skin EST 21-OCT-2002 ABV62925; ABV62925 standard; cDNA; 11 (first entry) 711. ВP

Human; skin; dermatological; vulnerary; antipsoriatic; antiseborrhaeic; immunosuppressive; antiinflammatory; cytostatic; SAGE; neurodermatitis; psoriasis; dermatitis; skin cancer; EST; expressed sequence tag; ss.

Homo sapiens.

WO200253774-A2

11-JUL-2002

20-DEC-2001; 2001WO-EP015179

03-JAN-2001; 2001DE-01000127

'n Conradt M, Hofmann

2002-590638/63

homeostasis and identifying In vitro identification of skin-expressed .g. skin cancer. cosmetic õ ed genes, useful pharmaceutical a agents for determining against

Disclosure; Page 45; 1345pp; German.

The invention relates to in vitro identification (M1) of genes expressed in the skin of humans or animals by subjecting a mixture of genetically encoded factors from skin, to serial analysis of gene expression (SAGE) so as to identify skin-expressed genes and quantify their expression. (M1) is useful for identifying genes involved in skin homeostasis; to determine skin homeostasis and to test agent (A) that maintains or promotes skin homeostasis or that can be used for treating skin disorders, specifically neurodermatitis, sumburn; psoriasis; scleroderma, other partitions are the standard of the sta ichthyosis; atopic dermatitis; acne; sebourhea; lupus erythematosus; rosacea; melanoma; basal cell carcinoma; and carcinoma or sarcoma of skin. The present sequence is that of a human expressed sequence tag (EST) of the invention carcinoma; and carcinoma or sarrooma of the that of a human expressed sequence tag scleroderma;

Disclosure; Page 224; 1345pp; German

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ABV68643;

ID ABV68643;

XX Human; ski

XX ABV68643;

YX W2002537;

PD 11-JUL-20;

XX O3-JAN-20;

YX O3-JAN-20;

YX O3-JAN-20;

YX O3-JAN-20;

YX D4 HENK ) HI

XX D4 HENK ) HI

XX D5 HOMEOSTABI

XX D7 HOMEOSTABI

XX D1 BClosur

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                                                                                                                 Query Match
Best Local S
Matches 10
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Best Local Similarity
Matches 10; Conserv
                                                                                                                                                                                                                                                                                                               The invention relates to in vitro identification (M1) of genes expressed in the skin of humans or animals by subjecting a mixture of genetically encoded factors from skin, to serial analysis of gene expression (SAGE) so as to identify skin-expressed genes and quantify their expression. (M1) is useful for identifying genes involved in skin homeostasis; to determine skin homeostasis and to test agent (A) that maintains or promotes skin homeostasis are that can be used for treating skin disorders, specifically neurodermatitis; sunburn, psoriasis; scleroderma; ichthyosis; atopic dermatities, cane; sebornhea; lupus erythematosus; rosacea; melanoma; basal cell carcinoma; and carcinoma or sarcoma of the skin. The present sequence is that of a human expressed sequence tag
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Disclosure; Page 204; 1345pp; German.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  In vitro identification of skin-expressed genes, useful for determining homeostasis and identifying cosmetic or pharmaceutical agents against
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Human; skin; dermatological; vulnerary; antipsoriatic; antiseborrhaeic; immunosuppressive; antiinflammatory; cytostatic; SAGE; neurodermatitis; psoriasis; dermatitis; skin cancer; EST; expressed sequence tag; ss.
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                                                                                                                                                                                                                               Sequence 11 BP; 3 A; 2 C; 3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       03-JAN-2001; 2001DE-01000127
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 20-DEC-2001; 2001WO-EP015179
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WO200253774-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human skin EST 6429
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      21-OCT-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (HENK ) HENKEL KGAA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1495 GACTTCAGCAG 1505
                                                       1489 AAGCCAGACTT 1499
                                                                                                                                                                                                                                                                                             of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    11
11
                                                                                                                                                Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    GACTACAGCAG 1
AAGCCAGTCTT 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     á
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                                                                                                                 Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       BP; 1 A; 3 C; 3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Conradt M,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    cDNA; 11
                                                                                                                                            31.3%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              31.3%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Hofmann K;
                                                                                                                                                                                                                               G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  0,
                                                                                                                 0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 G; 4 T; 0 U; 0 Other;
                                                                                                                                            Score 9.4;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Score 9.4;
Pred. No. 4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Mismatches
                                                                                                                 Mismatches
                                                                                                                                                                     9.4;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          DB 1;
                                                                                                                                                                     DB 1; Length 11;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  1;
                                                                                                              1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Length 11;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Indels
                                                                                                                 0;
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;-
                                                                                                              Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  0,
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RESULT 43

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RESULT 44
AAZ41704/c
ID AAZ4177
XX AAZ417
XX AAZ417
XX DT 20-MAR
DT 21-JAN
XX Organi
XX Amplif
KW detect
KW organo
                                                                                                                                                                                                                                                                                                                                                                        밁
                                                                                                                                                                                                                                                                             δ
                                                                                                                                                                                                                                                                                                          Query Match
Best Local Similarity
Matches 10; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                The invention relates to in vitro identification (M1) of genes expressed in the skin of humans or animals by subjecting a mixture of genetically encoded factors from skin, to serial analysis of gene expression (SAGE) so as to identify skin-expressed genes and quantify their expression. (M1) is useful for identifying genes involved in skin homeostasis; to determine skin homeostasis and to test agent (A) that maintains or promotes skin homeostasis or that can be used for treating skin disorders, specifically neurodermatitis; sunburn; psoriasis; scleroderma; ichthyosis; atopic dermatitis; acne; seborrhea; lupus erythematosus; rosacea; melanoma; basal cell carcinoma; and carcinoma or sarcoma of the skin. The present sequence is that of a human expressed sequence tag
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human; skin; dermatological; vulnerary; antipsoriatic; antiseborrhaeic; immunosuppressive; antiinflammatory; cytostatic; SAGE; neurodermatitis; psoriasis; dermatitis; skin cancer; EST; expressed sequence tag; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ABV69164 standard; cDNA; 11 BP
Amplification; polymerase chain reaction; PCR; microorganism; detection; pollutant; soil; food; agricultural chemical; polynorganochlorine; primer; ss.
                                                           Organic material detecting primer 65.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human skin EST
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ABV69164;
                                                                                          20-MAR-2003
21-JAN-2000
                                                                                                                                         AAZ41704;
                                                                                                                                                                                                                                                                                                                                                                         Sequence 11 BP; 7 A; 1 C; 3 G; 0 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Disclosure; Page 218; 1345pp; German.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         e.g. skin cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 2002-590638/63.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                03-JAN-2001; 2001DE-01000127.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   21-OCT-2002
                                                                                                                                                                    AAZ41704 standard; DNA; 12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       In vitro identification of skin-expressed genes, useful for determining homeostasis and identifying cosmetic or pharmaceutical agents against
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               20-DEC-2001; 2001WO-EP015179.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WO200253774-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (HENK ) HENKEL KGAA.
                                                                                                                                                                                                                                                                               1486 AAGAAGCCAGA 1496
                                                                                                                                                                                                                                                                                                                                                                                                        O.
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                                                                                                                                                                                                                                                                                                                                                                                                        the invention
                                                                                                                                                                                                                                                   AAGAAGCAAGA 11
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                                                                                                                                                                                                                                                                                                              Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (first entry)
                                                                                          (revised)
(first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Conradt
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      6950
                                                                                                                                                                                                                                                                                                                            31.3%;
90.9%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ₹,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Hofmann
                                                                                                                                                                       BP.
                                                                                                                                                                                                                                                                                                              <u>,,</u>
                                                                                                                                                                                                                                                                                                                            Score 9.4;
Pred. No. 4
                                                                                                                                                                                                                                                                                                               Mismatches
                                                                                                                                                                                                                                                                                                                               40;
                                                                                                                                                                                                                                                                                                                                           DB
                                                                                                                                                                                                                                                                                                                                            1.
                                                                                                                                                                                                                                                                                                              1;
                                                                                                                                                                                                                                                                                                                                           Length 11;
              polymer;
                                                                                                                                                                                                                                                                                                              0,
                                                                                                                                                                                                                                                                                                              Gaps
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RESULT 45
AAZ41488/c
ID AAZ414
XX AAZ414
XX II-JAN
DT 19-JAN
XX Microb
XX Microb
XX Microb
XX Fandom
XX III-JAN
XX III-JA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Ş
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 comprising (i) preparing many primers (P) with different probabilities of amplification and (ii) simultaneous polymerase chain reaction (PCR) of many different DNA using these primers. The method is used (i) to differentiate between different microorganisms in a mixed population and (ii) to determine presence/absence of an impurity (pollutant), or its concentration, in e.g. soil, foods, compost etc., typically metals, agricultural chemicals, polymers, organochlorine compounds etc. A particular use is monitoring composting of organic material.

Amplification with many primers produces a lot of information, so reliability of the test is improved, and many samples may be tested
                                                                                                                                                                                                                                                                                                                                         Microbe; detection; organic waste; arbitrarily primer PCR; random amplified polymorphic DNA; amplification; PCR prime
                                                                                                                                                                                                                                 JP11276176-A
                                                                                                                                                                                                                                                                                     Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                Microbe
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    19-JAN-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 12 BP; 1 A; 3 C; 4 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                quickly. AAZ41640-Z41855 represent the primers described in the invention. (Updated on 20-MAR-2003 to correct PR field.)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (SAOL )
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     31-MAR-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               30-MAR-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      21-OCT-1999.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Synthetic
                                                            31-MAR-1998;
                                                                                                                   31-MAR-1998;
                                                                                                                                                                          12-OCT-1999.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAZ41488 standard; DNA; 12.BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 This invention describes a novel method
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              16-MAR-1999;
        (SAOL ) SANYO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          1484 CCAAGAAGCCA 1494
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 10;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              polymerase chain reaction method, for differentiating between organisms and for detecting contaminants.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SANYO ELECTRIC CO LTD SOC TECHNO-INNOVATION
                                                                                                                                                                                                                                                                                                                                                                                                                             detection
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Page 18;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TECHNO-INNOVATION AGRIC FORESTY & FI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (first entry)
        ELECTRIC CO LTD
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              98JP-00087651.
99JP-00069694.
                                                            98JP-00087652.
                                                                                                                   98JP-00087652
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               99DE-01014461
                                                                                                                                                                                                                                                                                                                                                                                                                             in organic waste arbitrarily primed PCR primer #65
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        31.3%;
90.9%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ,
,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Score 9.4;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Pred. No. 43;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      for the amplification of DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     DB 1; Length 12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 <u>ب</u>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Indels
                                                                                                                                                                                                                                                                                                                                         primer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                in the method
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                of.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            of.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0
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Example; Page 9; 40pp; Japanese
                                                            Amplification of a DNA existence of a microbe.
                                                                fragment
                                                                 'n
                                                                 order to establish the
                                                                 state
                                                                 õ.
```

(NORI-) ZH NORIN SUISAN

SENTAN GIJUTSU SANGYO

A method has been developed for the amplification of a DNA fragment in which amplification is carried out on the DNA fragments of a number of different DNAs. The method comprises a PCR reaction repeatedly carrying out a heat-denaturing step, a primer annealing step and a polymerase extending step, to amplify the DNA fragments of a plural of different DNAs. The method can detect the existence of a microbe in organic waste. AAZ41424 to AAZ41639 represent PCR primers used in random amplified DNAs arbitrarily primed PCR, for the detection of microbes in organic waste

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Sequence 12 BP; 1 A;
3 C; 4
G; 4 T; 0 U; 0 Other;
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á Matches Query Match Local 1484 CCAAGAAGCCA 1494 l Similarity 10; Conserv Conservative 31.3%; Score 9.4; Pred. No. 4 Mismatches DB 1; Length 12; Indels 0 0

밁

CCAAGAGGCCA 1

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RESULT 46
AAC97839/c
JP2000270867-A
                                                                                                                   Synthetic
                                                                                                                                           28-FEB-2001
                                                                                                                                                   AAC97839
                                                                                                                                                          AAC97839
                                                                                                    03-OCT-2000
                                                                                                                           Primer; amplification; selective; ss.
                                                                                                                                   Primer used
                                                                                                                                                          standard;
                                                                                                                                   to illustrate DNA amplification method SEQ ID
                                                                                                                                          (first entry)
                                                                                                                                                          DNA;
                                                                                                                                                          ΒP
```

This invention relates to a method for amplifying a DNA fragment. The method comprises successive repetitions of heat-denaturing, annealing of a primer and an extending step using a DNA polymerase. The method makes use of a CDNA pool in which the primer is one primer or a pair of primer sets and has an amplification probability which allows it to amplify a DNA fragment from a limited number of the CDNAs among the DNA pool (where the limited number is in the range of 1 to 25). Also included in the invention are apparatus used for carrying out the method, a primer and a DNA polymerase and a kit used for amplifying a DNA fragment. The method can be used to amplify a limited number of cDNAs from a pool in which a wide variety of cDNAs are present. Oligonucleotides AAC97775 - AAC97990 represent primers used in an example illustrating the method of the

WPI; 2001-011047/02

Example 1; Page 8; 22pp; Japanese.

of a

DNA fragment

and

its

apparatus

(SAOL ) (NORI-)

SANYO ELECTRIC CO LTD. ZH NORIN SUISAN SENTAN

GIJUTSU SANGYO

19-MAR-1999; 19-MAR-1999;

99JP-00076844. 99JP-00076844.

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RESULT 47
ABIO2085/c
ID ABIO20
XX ABIO20
XX Oligon
XX SNP; s
KW peptid
KW Centra
XX SNP Homo s
XX Claim
XX Set of
PT design
PT design
PT design
PT design
PT design
CC This i
CC and c)
CC oligor
CC Tange
CC oligor
CC Tange
CC Claim
XX Claim

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                                                                                                                                                                                    Sequence 12 BP; 0 A; 1 C; 4 G; 7 T; 0 U; 0 Other;
                                                                                                                                                                                                                         was obtained in electronic format from W. ftp.wipo.int/pub/published_pot_sequences
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SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Set of oligonucleotides, useful for diagnosis and cell typing, designed to detect single-nucleotide polymorphisms and cytosine methylation status.
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                                                                                                                                 Oligonucleotide primer SEQ ID NO 353642 for detecting SNP TSC0048625.
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                                                           31-MAR-2000; 2000JP-00099482.
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                                                                                                                                                                             Synthetic
                                                                                                                                                                                                                                                    Microbial SSC-PCR primer SEQ ID NO 10
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Best Local 9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        The invention relates to identification of microbes, especially for obtaining information on the microbial spectrum in composted wastes and in the soil to which they are to be applied, and especially to give an indication of possible contaminants in the soil (such as mercury, arsenic, dioxins and environmental hormones) by the presence of microorganisms associated with them, comprising: (a) polymerase chain reaction on DNA of a mixture of microbes using a number of primers of different amplification probability (SSC-PCR); (b) electrophoresis of the DNA amplification fragments; (c) detecting the bands obtained on the electrophoresis image; (d) correcting errors in the image gradients; (e) measuring the position and intensity of the bands; (f) analyzing the results and creating a list of the band data; (g) searching the search. The present sequence is that of a PCR primer, useful to the invent
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Microbe identification comprises comparing results of reference database for determining microbial spectrum
                              Diagnosis of cancer in potentially neoplastic samples - by comparing the level of transcription between RNA transcripts in two tissue samples, useful for providing an extensive profile of gene expression in rat embryo fibroblast (REF) cells.
                                                                                                                                                                                                                                                                                                                                                                                                                      02-JUL-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       p53; serial analysis of gene expression; SAGE tag; cancer; neoplastic; rat embryo fibroblast; REF; tumour suppressor; cell cycle control;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          p53 serial analysis of gene expression tag #45
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                                                                                                                                                                                  WPI; 1999-106079/09.
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                                                                                                                                                                                                                                            Galella EA,
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Pred. No.
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               19-JUN-1998;
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19-JUN-1998;
19-JUN-1998;
19-JUN-1998;
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                                                                                                                                  19-JUN-1998
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                                                                                                                                                                                                                                                                                                                                                                                                                               cytotoxic T-lymphocyte; tumour antigen;
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19-JUN-1998;
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98US-0090036P
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0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                              immunotherapy; anticancer; ss
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RESULT 53
AAZ83241/c
ID AAZ832
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AC AAZ832
XX

AAZ83241 standard; DNA; 10 BP

AAZ83241;

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Sequence 10

BP; 1

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2 C; 4 G; 3 T; 0

U; 0 Other; DB 1;

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other transcripts correspond to novel genes. Antigen-presenting cells (APC) associated costimulatory factors play an important role in the cativation of the cytotoxic immune response, particularly against tumour cells. Tumour antigen presentation via the MHC (major histocompatibility complex) and subsequent recognition by T-cell receptors is alone insufficient to activate a robust cytotoxic immune response that can lyse the tumour cells, immunostimulatory cofactors also being required for cefficient activation of cytotoxic T-lymphocytes (CTLs). Nucleic acid sequences identified using the SAGE tags have several potential uses. The pay be used in vaccines to induce an immune response, particularly against a tumour antigen; to modulate the genotype of an APC; to screen for agents that modulate expression of differentially expressed genes in an APC; and as hybridisation probes/amplification primers for the cypressed genes, or of their encoded proteins, can be used to identify cells as belonging to the monocyte lineage. Cells containing these genes can be used in active immunotherapy (or to stimulate production of a propulation of antigen-specific effector cells) and vectors containing these genes compulation of antigen-specific effectors ensures adequate antigen and presentation of co-stimulatory factors ensures adequate antigen the gresentation of co-stimulatory signals, migration to Tcell-rich sites, recruitment of immune effector cells
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       19-JUN-1998;
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19-JUN-1998;
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19-JUN-1998;
19-JUN-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         with monocytes. Some of the transcripts correspond to known genes or (expressed sequence tags) which were previously unknown to be preferentially or differentially expressed in dendritic cells, while
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequences AAZ77573-Z79709 represent SAGE (serial analysis of gene expression) tags used to identify mRNA transcripts encoding immunostimulatory cofactor proteins which are preferentially or differentially expressed in monocyte-derived dendritic cells compared
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Claim 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       cells, useful in gene vaccines against cancer.
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(ROBE/)
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ROBERTS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           polynucleotides differentially expressed in antigen-presenting
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98US-0090048P.
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RESULT 54
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                                                                                                                                                                                                                                                               treatment of breast cancer, particularly where metastatic. Diagnosis is by standard immunoassays or hybridisation/amplification reactions. Compounds that modulate expression of the transcripts are potentially useful for treatment of (metastatic) breast cancer, while promoters from the transcripts are used to direct expression, in selected cell types, of e.g. therapeutic genes (also ribozymes or antisense sequences), particularly an antigen-encoding sequence for use in gene or cell-based vaccines. Polypeptides encoded by the transcripts are also useful in vaccines; for diagnosing breast cancer and for raising specific antibodies (Ab). Ab are used to detect the polypeptides or as therapeutic agents. Host cells that produce the polypeptides can be used to expand and isolate populations of educated, antigen-specific immune effecter cells, e.g. cytotoxic T lymphocytes, and these used for adoptive
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAZ80767 to AAZ83941 represent tags corresponding to distinct transcripts that are preferentially transcribed in the metastatic breast tumour tissue (i.e. are upregulated in metastatic breast tumour cells). AAZ83942 to AAZ86677 represent tags corresponding to distinct transcripts that are preferentially transcribed in the primary or non-metastatic breast tumour tissue (i.e. are downregulated in metastatic breast tumour cells). These transcripts can be used for diagnosis, prognosis, monitoring and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human; metastatic breast tumour tissue; breast cancer; tag; primer; non-metastatic breast tumour tissue; gene therapy; anticancer; antimetastatic; vaccine; diagnosis; ss.
              AAZ83053 standard; DNA; 10 BP
                                                                                                                                                                                                                      Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Isolated polynucleotides differentially expressed between metastatic non-metastatic breast cancer cells, useful for diagnosis, prevention
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SHANKARA S.
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19-JUN-1998;
                                                                                                                                                                              e.g. therapeutic genes (also ribozymes or antisense sequences), particularly an antigen-encoding sequence for use in gene or cell-based vaccines. Polyapptides encoded by the transcripts are also useful in vaccines; for diagnosing breast cancer and for raising specific antibodies (Ab). Ab are used to detect the polyapptides or as therapeutic agents. Host cells that produce the polyapptides can be used to expand and isolate populations of educated, antigen-specific immune effecter cells, e.g. cytotoxic T lymphocytes, and these used for adoptive
                                                                                                                                                                                                                                                                                                                 to AAZ86677 represent tags corresponding to distinct transcripts that are preferentially transcribed in the primary or non-metastatic breast tumour tissue (i.e. are downregulated in metastatic breast tumour cells). These transcripts can be used for diagnosis, prognosis, monitoring and treatment of breast cancer, particularly where metastatic. Diagnosis is by standard immunoassays or hybridisation/amplification reactions. Compounds that modulate expression of the transcripts are potentially useful for treatment of (metastatic) breast cancer, while promoters from the transcripts are used to direct expression, in selected cell types, of
                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAZ80767 to AAZ83941 represent tags corresponding to distinct transcripts that are preferentially transcribed in the metastatic breast tumour tissue (i.e. are upregulated in metastatic breast tumour cells). AAZ83942 to AAZ86677 represent tags corresponding to distinct transcripts that are
                                                                                                                                 Sequence 10 BP; 1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Isolated polynuclectides differentially expressed between metastatic non-metastatic breast cancer cells, useful for diagnosis, prevention
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(ROBE/)
(SHAN/)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human; metastatic breast tumour tissue; breast cancer; tag; primer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Metastatic breast tumour cell upregulated
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ROBERTS B L.
SHANKARA S.
                                                              Similarity 9; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          of cancer.
                                                                  Conservative
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98US-0089997P.
98US-0090039P.
98US-0090040P.
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RESULT 55 AAZ86659

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Matches 9
                                                                                                                                                                                                                                                                                                     e.g. therapeutic genes (also ribozymes or antisense sequences), particularly an antigen-encoding sequence for use in gene or cell-based vaccines. Polypeptides encoded by the transcripts are also useful in vaccines; for diagnosing breast cancer and for raising specific antibodies (Ab). Ab are used to detect the polypeptides or as therapeutic agents. Host cells that produce the polypeptides can be used to expand and isolate populations of educated, antigen-specific immune effecter cells, e.g. cytotoxic T lymphocytes, and these used for adoptive
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAZ80767 to AAZ83941 represent tags corresponding to distinct transcripts that are preferentially transcribed in the metastatic breast tumour tissue (i.e. are upregulated in metastatic breast tumour cells). AAZ83942 to AAZ86677 represent tags corresponding to distinct transcripts that are preferentially transcribed in the primary or non-metastatic breast tumour cells). These transcripts can be used for diagnosis, prognosis, monitoring and treatment of breast cancer, particularly where metastatic. Diagnosis is by standard immunoassays or hybridisation/emplification reactions. Compounds that modulate expression of the transcripts are potentially useful for treatment of (metastatic) breast cancer, while promoters from the transcripts are used to direct expression, in selected cell types, of
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(ROBE/)
(SHAN/)
                                                                                                                                                                                                                             Sequence 10 BP; 6 A; 2 C; 2 G; 0 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             treatment of cancer.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       07-APR-2000
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19-JUN-1998;
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                                                                                                                                        Local Similarity
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ROBERTS B L.
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                                                      AAGAAGCCA 1494
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                                                                                                                Conservative
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Pred. No.
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RESULT 57
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AAF70127 standard; DNA; 10

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                                                                                                  a cell expressing the target epitope with cytotoxic T cells generated against the cell to identify DNA clones expressing the target epitope. The method may also comprise providing a cytotoxic T cell specific for a centrifying tumour specific T cell. The methods are useful for ceativity of the cytotoxic T cell. The methods are useful for centrifying tumour specific target epitopes and antigens which are useful in immunogenic compositions or vaccines to induce the regression of cumours, cancers or infections in mammals. The genes expressed in a panel contingenic cell line are used to generate HLA restricted cytotoxic T cells which are evaluated for activity against tumour cells. The method is useful to identify potential antigens expressed not only by the pathogen but also by the host cells whose gene expression is altered as a result of infection. The differential gene expression strategies can be applied to identify immunogenic molecules of cells infected with virus,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAA15239-50 represent arbitrary primers which are used for modified differential display of tumour antigens, in the method of the invention. The specification describes a method for identifying a target epitope. The method comprises screening the products of an expression library from a cell expressing the target epitope with cytotoxic T cells generated
    Sequence 10 BP; 3 A; 3 C; 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Synthetic.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Disclosure; Page 68; 132pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Novel method of identifying target epitopes or antigens specific for human tumors, cancers and infected cells involving screening express
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                10-NOV-1998;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    library products of a cell expressing the target epitope.
                                                                                  or mycobacterium
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human; TNFRSF11B; osteoclastogenesis inhibitory factor; single nucleotide polymorphism; SNP; osteoclast recruitment; osteoclast function; osteoporosis; metastatic bone disease; Paget's disease; rheumatoid arthritis; periodontal bone disease;
                                                                                       nor previously assigned open reading serial analysis of gene expression; a
                                                                                                   Yeast; Saccharomyces cerevisiae; characterisation; cell cycle;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human Osteoclastogenesis Inhibitory Factor nucleotides, comprising single nucleotide polymorphisms, useful for studying e.g. osteoporosis, Paget's disease and rheumatoid arthritis.
                                                 Saccharomyces cerevisiae
                                                                                                                                             Yeast NORF
                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 10
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21-DEC-2000
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                                                                                                                                           SAGE tag oligonucleotide SEQ ID NO:1748.
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cc antifungal drugs comprising: (a) contacting a test substance with a yeast cell; and (b) monitoring expression of a NORF gene whose expression of cvaries as in M1, where a test substance which modifies the expression of the yeast gene is a candidate antifungal drug; (3) a method (M3) for comprising contacting human DNA with a probe which comprises at least 10 contiguous nucleotides of a NORF gene whose expression varies as in M1; candidate of the contiguous nucleotides of a NORF gene whose expression varies as in M1; candidate of drugs having a characteristic effect on gene expression in a collass of drugs having a characteristic effect on gene expression in a collass of drugs having a characteristic effect on gene expression in a collass of drugs and it is affected by the class of drugs. The NORF gene whose expression is affected by the class of drugs. The NORF genes may be used to identify candidate drugs which affect the cell cycle. The cepresent sace tags used in the exemplification of the present invention. CC AAF33267 represent linkers and PCR primers used in the SAGE method, in the exemplification of the present invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   The present invention describes an isolated DNA molecule comprising a coding sequence of a yeast gene selected from a group of 745 NORF (not previously assigned open reading frame; or nonannotated ORF) genes comprising a SAGE (serial analysis of gene expression) tag. Also described are: (1) a method (M1) of using NORF genes to affect the cell cycle comprising administering a NORF gene whose expression varies by at least 10% between any two phases of the cell cycle selected from log phase, S phase and G2/M; (2) a method (M2) for screening candidate antifungal drugs comprising: (a) contacting a test substance with a yeas
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Yeast gene coding sequences comprising NORF genes with serial analysis gene expression (SAGE) tags, useful for studying, monitoring and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        14-JUN-2000; 2000WO-US016223
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                phases of the cell cycle.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 62; 419pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Vogelstein B,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         9908-00335032
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Sequence 10 BP; 2 A; 2 C; 3 G; 3 T; 0 U; 0 Other;

Matches Query Match 1498 TTCAGCAGC 1506 l Similarity 9; Conserv Conservative 30.0%; 100.0%; Score 9; Pred. No. Mismatches DB . ۲, Length 10; 0 0; Gaps 0

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10

TTCAGCAGC

AAF39472 standard; DNA; 10 ВP

23-MAR-2001 gene (first entry) SAGE tag oligonucleotide SEQ ID NO:6211.

RESULT 59
AAF39472/c
ID AAF394
XX AAF394
XX AAF394
XX Yeast
XX Yeast
XX Yeast
XX Yeast
XX Yeast
XX Yeast
XX Inker
XX Saccha
XX Saccha Yeast; Saccharomyces cerevisiae; characterisation; cell cycle; NORF; nor previously assigned open reading frame; nonannotated ORF; SAGE; serial analysis of gene expression; antifungal; tag; identification; PCR primer;

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RESULT 60
AAF38820
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CC comprising a SAGE (serial analysis of gene expression) tag. Also codescribed are: (1) a method (M1) of using NORF genes to affect the cell cycle comprising administering a NORF gene whose expression varies by at CC least 10% between any two phases of the cell cycle selected from log CC phase, S phase and G2/M; (2) a method (M2) for screening candidate CC enliftungal drugs comprising: (a) contacting a test substance with a yeast CC cell; and (b) monitoring expression of a NORF gene whose expression of CC tries as in M1, where a test substance which modifies the expression of CC tries as in M1, where a test substance which modifies the expression of CC comprising contacting human genes which are involved in cell cycle progression of CC contiguous nucleotides of a NORF gene whose expression varies as in M1; CC contiguous nucleotides of a NORF gene whose expression varies as in M1; CC contiguous nucleotides of a NORF gene whose expression in a CC class of drugs having a characteristic effect on gene expression in a CC class of drugs having a characteristic effect on gene expression in a CC comprising contacting a yeast cell with a candidate drug and contacting expression in the yeast cell of at least 1 NORF gene whose expression is affected by the class of drugs. The NORF gene whose used CC expression is affected by the class of the cell cycle, the differentially contacting the contacting the contacting the contacting tries of the cell cycle.
                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
Best Local Similarity
Matches 9; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              expressed genes may be used as markers of phases of the cell cycle. The methods may be used to identify candidate drugs which affect the cell cycle and for identification of antifungal drugs. AAF33268 to AAF44064 represent SAGE tags used in the exemplification of the present invention AAF33362 to AAF33267 represent linkers and PCR primers used in the SAGE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 The present invention describes an isolated DNA molecule comprising coding sequence of a yeast gene selected from a group of 745 NORF (repreviously assigned open reading frame; or nonannotated ORF) genes comprising a SAGE (serial analysis of gene expression) tag. Also
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Yeast gene coding sequences comprising NORF genes with serial analysis gene expression (SAGE) tags, useful for studying, monitoring and affecting phases of the cell cycle.
Yeast; Saccharomyces cerevisiae; characterisation; cell cycle; NORF; nor previously assigned open reading frame; nonamotated ORF; SAGS; serial analysis of gene expression; antifungal; tag; identification; linker; PCR primer; ds.
                                                                                                                                                                                                                                     AAF38820 standard; DNA; 10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 10 BP; 1 A; 2 C; 2 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Example; Page 221; 419pp; English.
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                                                                                                         Yeast
                                                                                                                                                    23-MAR-2001
                                                                                                                                                                                             AAF38820
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                                                                                                                                                                                                                                                                                                                                                                                        1483 ACCAAGAAG 1491
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                                                                                                       gene
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               the exemplification of the present invention
                                                                                                                                                                                                                                                                                                                                                                                                                                   Conservative
                                                                                                                                                  (first entry)
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                                                                                                         SAGE
                                                                                                     tag oligonucleotide
                                                                                                                                                                                                                                                                                                                                                                                                                                                       100.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              30.0%;
                                                                                                                                                                                                                                     ВP
                                                                                                                                                                                                                                                                                                                                                                                                                                Score 9; DB 1; Pred. No. 43; 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                   0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Kinzler K;
                                                                                                                                                                                                                                                                                                                                                                                                                                                       DB 1;
                                                                                                         SEQ
                                                                                                         IJ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Length 10,
                                                                                                                                                                                                                                                                                                                                                                                                                                0;
                                                                                                         NO:5559
                                                                                                                                                                                                                                                                                                                                                                                                                                   Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                0;
                                                                                                                                                                                                                                                                                                                                                                                                                                   Gaps
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The present invention describes an isolated DNA molecule comprising a CC coding sequence of a yeast gene selected from a group of 745 NORF (not previously assigned open reading frame; or nonannotated ORF) genes CC comprising a SAGE (serial analysis of gene expression) tag. Also CC described are: (1) a method (M1) of using NORF genes to affect the cell cycle comprising administering a NORF gene whose expression varies by at CC least 10% between any two phases of the cell cycle selected from log phase, S phase and G2/M; (2) a method (M2) for screening candidate antifungal drugs comprising: (a) contacting a test substance with a yeast CC cell; and (b) monitoring expression of a NORF gene whose expression of the yeast gene is a candidate antifungal drug; (3) a method (M3) for comprising contacting human DNA with a probe which comprises at least 10 comprising contacting human DNA with a probe which comprises at least 10 comprising contacting human DNA with a probe which comprises at least 10 cc candidate of a NORF gene whose expression in CC cand (4) a method (M4) for identifying a candidate drug as a member of a cyeast cell comprising contacting a yeast cell with a candidate drug and monitoring expression in the yeast cell of at least 1 NORF gene whose expression in a cyeast cell of the least 1 NORF gene whose expression in a cyeast cell of the cell cycle. The method (M3) be used to identify candidate drugs. The NORF genes may be used to study, monitor and affect phases of the cell cycle. The cycle and for identification of antifungal drugs. AAF33268 to AAF44064 crepressent SAGE tags used in the exemplification of the present invention. AAF33262 to AAF33267 represent linkers and PCR primers used in the SAGE method, in the exemplification of the present invention.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Yeast gene coding sequences comprising NORF genes with serial analysis gene expression (SAGE) tags, useful for studying, monitoring and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WPI; 2001-061874/07.
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O.f

Sequence 10 BP; 6 A; 2 C; 2 G; 0 T; 0 U; 0 Other;

B Ş Matches Query Match Best Local Similarity 1486 AAGAAGCCA 1494 9; Conservative ý 30.0%; Score 9; 100.0%; Pred. No. 0 Mismatches DB 1; Length 10; <u>.</u> Indels ٥,

0,

AAF41980
ID AAF4
XX AC AAF4
XX AC AAF4
XX Yeas
XX Yeas
XX Yeas AAF41980; Yeast NORF gene SAGE tag oligonucleotide SEQ ID NO:8719 23-MAR-2001 (first entry)

RESULT 61

AAF41980 standard;

DNA;

10

Yeast; Saccharomyces cerevisiae; characterisation; cell cycle; NORF;

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RESULT 62
AAF38821
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AhF33267 to AhF33267 represent linkers and PCR primers used in the SAGE method in the SAGE
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XXXXXX
                                                                                                                                                                                                                                                                                                 Query Match
Best Local Similarity
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                23-MAR-2001
                                                        AAF38821;
                                                                                                                                                                                                                                                                                                                                                            Sequence 10
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                                                                                            AAF38821 standard; DNA; 10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Velculescu V,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          14-JUN-2000; 2000WO-US016223
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                                                                                                                                                                                                                                                                                                                                                                                                    in the
                                                                                                                                                                                                                                         AGAAGCCAG 1495
                                                                                                                                                                                                                                                                                                                                                            BP; 4 A;
                                                                                                                                                                                                                                                                               Conservative
              (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                    exemplification of the present invention
                                                                                                                                                                                                 10
                                                                                                                                                                                                                                                                                               30.0%;
                                                                                                                                                                                                                                                                                                                                                            2 C; 4
                                                                                                                                                                                                                                                                               0
                                                                                                                                                                                                                                                                                                 Score 9;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                            G; 0 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                 Mismatches
                                                                                                                                                                                                                                                                                                 DB 1;
b. 43;
                                                                                                                                                                                                                                                                                                                    Length 10;
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                                                                                                                                                                                                                                                                               Indels
                                                                                                                                                                                                                                                                               0,
                                                                                                                                                                                                                                                                               Gaps
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Yeast; Saccharomyces cerevisiae; characterisation; cell cycle; NORF; nor previously assigned open reading frame; nonannotated ORF; SAGE;
                                                                                                                                                                         Yeast NORF gene SAGE tag oligonucleotide
                                                                                                                                                                          SEQ
                                                                                                                                                                          IJ
                                                                                                                                                                          NO:5560
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nor previously assigned open reading frame; nonannotated ORF; SAGE; serial analysis of gene expression; antifungal; tag; identification PCR primer; ds.

Saccharomyces cerevisiae

14-JUN-2000; 2000WO-US016223

16-JUN-1999; 99'08-00335032

(UYJO ) UNIV JOHNS HOPKINS

Velculescu V, Vogelstein B, Kinzler

affecting phases of the cell cycle. Yeast gene coding sequences comprising NORF genes with serial analysis gene expression (SAGE) tags, useful for studying, monitoring and of.

Example; Page 198; 419pp; English.

The present invention describes an isolated DNA molecule comprising a coding sequence of a yeast gene selected from a group of 745 NORF (not previously assigned open reading frame, or nonannotated ORF) genes comprising a SAGE (serial analysis of gene expression) tag. Also comprising administering a NORF genes to affect the cell cycle comprising administering a NORF gene whose expression varies by at cell east 10% between any two phases of the cell cycle selected from log cantifungal drugs comprising; (a) contacting a test substance with a yeast cell; and (b) monitoring expression of a NORF gene whose expression of the yeast gene is a candidate antifungal drug; (3) a method (M3) for contacting human DNA with a probe which comprises at last 10 contiguous nucleotides of a NORF gene whose expression of contiguous nucleotides of a NORF gene whose expression of contiguous nucleotides of a NORF gene whose expression of a yeast cell comprising contacting human DNA with a probe which comprises at least 10 contacting a characteristic effect on gene expression in a yeast cell comprising contacting a yeast cell with a candidate drug and contacting a yeast cell with a candidate drug and yeast cell comprising contacting a yeast cell with a candidate drug and contacting a yeast cell with a candidate drug and contacting a yeast cell of at least 1 NORF gene whose expression in a grees of drugs having a characteristic effect on gene expression in a contacting a yeast cell with a candidate drug and contacting a yeast cell of at least 1 NORF genes may be used to study, monitor and affect phases of the cell cycle, the differentially expressed genes may be used as markers of phases of the cell cycle. The Contact of the cell cycle and foor identification of antifungal drugs. ARR33268 to ARR4064 contact of the present invention.

Contact of the present invention. 'n the exemplification of the present invention

Sequence 10 ₿₽; 5 A; 2 C; 2 G; 1 T; 0 U; 0 Other;

В S Query Match Best Local S Matches S 1486 \_ l Similarity 9; Conserv AAGAAGCCA 1494 Conservative 100.0%; 30.0%; 0; Score 9; 1 Pred. No. Mismatches DB 43; 1 Length 10; Indels 0

Gaps

0

RESULT 63
AAF39528
ID AAF39
XX
AC AAF39 AAF39528 standard; DNA; 10

AAF39528

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                                                                                                                                                cc antifungal drugs comprising: (a) contacting a test substance with a yeast ccell; and (b) monitoring expression of a NORF gene whose expression of the yeast gene is a candidate antifungal drug; (3) a method (M3) for CC the yeast gene is a candidate antifungal drug; (3) a method (M3) for CC contiguous nucleotides of a NORF gene whose expression of CC contiguous nucleotides of a NORF gene whose expression varies as in M1; CC and (4) a method (M4) for identifying a candidate drug as a member of a CC class of drugs having a characteristic effect on gene expression in a CC yeast cell comprising contacting a yeast cell with a candidate drug and CC monitoring expression in the yeast cell of at least 1 NORF gene whose cexpression is affected by the class of drugs. The NORF genes may be used to study, monitor and affect phases of the cell cycle, the differentially cycle and for identification of antifungs. AAF33268 to AAF33267 represent in the exemplification of the present intention. CC AAF33262 to AAF33267 represent linkers and PCR primers used in the SAGE contaction of the present invention.
                                                                                       Query Match
Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       previously assigned open reading frame; or nonannotated ORF) genes comprising a SAGE (serial analysis of gene expression) tag. Also described are: (1) a method (M1) of using NORF genes to affect the cell cycle comprising administering a NORF gene whose expression varies by at least 10% between any two phases of the cell cycle selected from log phase, S phase and G2/M; (2) a method (M2) for screening candidate
                                                                                                                                              Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          The present invention describes an isolated DNA molecule comprising coding sequence of a yeast gene selected from a group of 745 NORF ()
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Yeast gene coding sequences comprising NORF genes with serial analysis of gene expression (SAGE) tags, useful for studying, monitoring and affecting phases of the cell cycle.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Velculescu V, Vogelstein B,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Yeast; Saccharomyces cerevisiae; characterisation; cell cycle; NORF; nor previously assigned open reading frame; nonannotated ORF; SAGE; serial analysis of gene expression; antifungal; tag; identification;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            16-JUN-1999;
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                                                                        9
                                                                                       Similarity
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                                 AGAAGCCAG 1495
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                                                                        Conservative
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                                                                                       30.0%;
                                                                                                                                            2 C; 3 G; 1 T; 0 U; 0 Other;
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                                                                      0
                                                                                       Score 9;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Kinzler K;
                                                                        Mismatches
                                                                                                         DB 1;
                                                                                                       Length 10;
                                                                    0;
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                                                                      Indels
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RESULT 64 AAF38557

8 S

1489

AAGCCAGAC 1497 AAGCCAGAC

10

Query Match Best Local S Matches 9

Similarity 9; Conserv

Conservative

0

Sequence 10

BP; 5 A; 3 C; 2 G; 0 T; 0 U; 0 Other;

30.0%; Score 9; 100.0%; Pred. No.

DB 1;

Length 10

0

Gaps

0;

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CC previously assigned open reading frame; or nonamotated ORP) genes (CC comprising a SAGE (serial analysis of gene expression) tag. Also (CC described are: (1) a method (M1) of using NORF genes to affect the cell (CC cycle comprising administering a NORF gene whose expression varies by at (CC least 10% between any two phases of the cell cycle selected from log (CC phase, S phase and G2/M; (2) a method (M2) for screening candidate (CC antifungal drugs comprising: (a) contacting a test substance with a yeast (cell; and (b) monitoring expression of a NORF gene whose expression of (CC varies as in M1, where a test substance which modifies the expression of (CC contiguous mucleotides of a NORF gene whose expression of (CC contiguous mucleotides of a NORF gene which comprises at least 10 (CC contiguous mucleotides of a NORF gene whose expression of (CC contiguous mucleotides of a NORF gene whose expression varies as in M1; (CC and (4) a method (M4) for identifying a candidate drug as a member of a (CC class of drugs having a characteristic effect on gene expression in a cyeast cell with a candidate drug and (CC monitoring expression in the yeast cell with a candidate drug and (CC expressed genes may be used as markers of phases of the cell cycle. The (CC cycle and for identify candidate drugs, MAF33268 to AAF33267 represent 1inkers and PCR primers used in the SAGE (CC AAF33262 to AAF33267 represent linkers and PCR primers used in the SAGE (CC method, in the exemplification of the present invention (CC AAF33262 to AAF33267 represent linkers and PCR primers used in the SAGE (CC method, in the exemplification of the present invention (CC AAF33267 to AAF3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Yeast gene coding sequences comprising NORF genes with serial analysis gene expression (SAGE) tags, useful for studying, monitoring and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Yeast; Saccharomyces cerevisiae; characterisation; cell cycle; NORF; nor previously assigned open reading frame; nonamotated ORF; SAGE; serial analysis of gene expression; antifungal; tag; identification; linker; PCR primer; ds.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               The present invention describes an isolated DNA molecule comprising a coding sequence of a yeast gene selected from a group of 745 NORF (not
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Example; Page 189; 419pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                affecting phases of the cell cycle.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WPI; 2001-061874/07.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Velculescu V,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      16-JUN-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           14-JUN-2000; 2000WO-US016223
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Saccharomyces cerevisiae.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Yeast NORF gene SAGE tag oligonucleotide SEQ ID NO:5296
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     23-MAR-2001
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (UYJO ) UNIV JOHNS HOPKINS.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        9908-00335032
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RESULT 65
AAD31785
                                                                       Query Match
Best Local Similarity
"---hes 9; Conserva
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                                                                                                                                                                                                                                                                                                                                                                                            The present invention relates to novel methods for the identification of antigens recognised by cytotoxic T cells (CTLs) and specific for human tumours, cancers and infected cells. The method involves screening the products of an expression library generated from DNA/RNA of a cell cell to identify DNA clones expressing target epitope or providing cytotoxic T cells specific for a gene product differentially expressed by a cell and measuring the cross-reactivity of the cytotoxic T cells for cells expressing a target epitope in which the target epitope is cidentified as a gene product inducing cytotoxic T cells. The method is constituted for identifying a target epitope or antigen specific for a tumour cell. The target epitope is also useful for identifying target antigens in other target cells against which it is desirable to induce cell-conditated immunity. The antigen identified by the method is useful in immunogenic compositions and vaccine preparations to induce the invention calso relates to vaccinia viral vectors which are useful for treating the compositions and varial vectors which are useful for treating the compositions and varial vectors which are useful for treating the compositions and varial vectors which are useful for treating the compositions and varial vectors which are useful for treating the compositions and varial vectors which are useful for treating the compositions and varial vectors which are useful for treating the compositions and varial vectors which are useful for treating the compositions and varial vectors which are useful for treating the compositions and varial vectors which are useful for treating the compositions and varial vectors which are useful for treating the compositions and varial vectors which are useful for treating the compositions and the c
                                                                                                                                                                                                                                                                                                                          related to vaccinia the response tumour-bearing mammals, including humans to generate immune response against the tumour cells. They are also useful for immunising or vaccinating tumour-free subjects to prevent tumour formation. The present
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAD31785 standard; DNA; 10 BP
                                                                                                                                                                                                                                                  vaccinating tumour-free subjects to prevent tumour formation. The present DNA sequence is an arbitrary primer which is used for modified differential display of genes encoding potential tumour immunogens. This primer is used in the exemplification of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Representational Difference Analysis method for identification of antigens recognized by cytotoxic T cells and specific for human tumors,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Unidentified
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      MR 7 arbitrary primer used for modified differential display.
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                                                                                                                                                                                                  Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Example 4; Page 19; 54pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         comprises improved selection of genes encoding target antigens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WPI; 2002-239252/29.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    02-APR-2001; 2001US-00822250.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Cytotoxic T cell;
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                                             1493 CAGACTTCA 1501
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CAGACTTCA 10
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                                                                                               Conservative
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A;
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response; cytostatic; primer; ss.
                                                                                                                     30.0%;
                                                                                                                                                                                                  3 C; 1 G; 3 T; 0 U; 0 Other;
                                                                                               0
                                                                                                                     Score 9; DB 1
Pred. No. 43;
                                                                                               Mismatches
                                                                                                                                             DB 1;
                                                                                                                                             Length 10;
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                                                                                               Indels
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                                                                                               Gaps
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RESULT 66 ABQ87208/c

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ABV69313/c
ABV69313/c
ID ABV693
XX
AC ABV693
XY
C 21-OCT
XX
DT 21-OCT
XX
Human
XX
M immuno
XW immuno
XW jsoria
XX
OS Homo s
XX
PN WO2002
XX
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Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                      expression. (A) comprises protein or making to the comprises protein or making and/or stress; determining useful for: identifying markers of skin ageing and/or stress; and identifying or determining the effects of pharmaceutical or cosmetic agents for control of skin ageing. The present sequence is one of a group of human skin ageing/stress related expressed sequence tags (ABQ86246-ABQ87680) of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        The invention relates to identifying (M1) genes in vitro that, in humans or animals, are important for skin ageing and/or skin stress by serial analysis of gene expression between mixtures of transcribed and optionally translated, genetically encoded factors (A) obtained from young and aged skin, to identify that genes that show strong differential expression. (A) comprises protein or mRNAs or their fragments. (M1) is
                                                                       Human; skin; dermatological; vulnerary; antipsoriatic; antiseborrhaeic; immunosuppressive; antiinflammatory; cytostatic; SAGE; neurodermatitis; psoriasis; dermatitis; skin cancer; EST; expressed sequence tag; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human; skin ageing; skin stress; EST;
                                                                                                                                     Human skin EST
                                                                                                                                                                   21-OCT-2002
                                                                                                                                                                                                ABV69313;
                                                                                                                                                                                                                            ABV69313 standard; cDNA; 11
                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 11 BP; 1 A; 2 C; 4 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Claim 8; Page 77; 325pp; German
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Identifying genes involved in skin stress and aging, useful e.g. in screening for cosmetic or therapeutic agents, based on differential gene
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WPI;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 03-JAN-2001; 2001DE-01000121
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WO200253773-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Homo sapiens.
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                                             Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (HENK ) HENKEL KGAA.
                                                                                                                                                                                                                                                                                                                           1490 AGCCAGACT 1498
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                                                                                                                                                                                                                                                                                                     11 AGCCAGACT 3
                                                                                                                                                                                                                                                                                                                                                             Similarity 9; Conserv
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                                                                                                                                                                                                                                                                                                                                                                Conservative
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                                                                                                                                                                   (first entry)
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                                                                                                                                     7099
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                                                                                                                                                                                                                                                                                                                                                                                              30.0%;
                                                                                                                                                                                                                                                                                                                                                                            100.0%;
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                                                                                                                                                                                                                              ВP
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                                                                                                                                                                                                                                                                                                                                                                Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   expressed sequence tag; ss
                                                                                                                                                                                                                                                                                                                                                                            DB 1;
o. 47;
                                                                                                                                                                                                                                                                                                                                                                                         Length 11;
                                                                                                                                                                                                                                                                                                                                                                0,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 963
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0

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RESULT 68
ABV71907/c
ID ABV71
XX ABV71
XX ABV71
XX ABV71
XX ABV71
XX 21-OC
XX Humar
XX Humar
XX Humar
XX Homo

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Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    The invention relates to in vitro identification (M1) of genes expressed in the skin of humans or animals by subjecting a mixture of genetically encoded factors from skin, to serial analysis of gene expression (SAGE) so as to identify skin-expressed genes and quantify their expression (M1) is useful for identifying genes involved in skin homeostasis; to determine skin homeostasis and to test agent (A) that maintains or promotes skin homeostasis or that can be used for treating skin disorders, specifically neurodermatitis; sunburn; psoriasis; scleroderma; ichthyosis; atopic dermatitis; acne; seborrhe; lupus erythematosus; rosacea; melanoma; basal cell carcinoma; and carcinoma or sarcoma of the skin. The present sequence is that of a human expressed sequence tag (EST) of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Human; skin; dermatological; vulnerary; antipsoriatic; antiseborrhaeic; immunosuppressive; antiinflammatory; cytostatic; SAGE; neurodermatitis; psoriasis; dermatitis; skin cancer; EST; expressed sequence tag; ss.
                                                                WPI; 2002-590638/63
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human skin EST 9693
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 In vitro identification of skin-expressed genes, useful for determining homeostasis and identifying cosmetic or pharmaceutical agents against
In vitro identification of skin-expressed genes, useful for determining
                                                                                                                       Petersohn D,
                                                                                                                                                                                                                                    03-JAN-2001; 2001DE-01000127
                                                                                                                                                                                                                                                                                              20-DEC-2001; 2001WO-EP015179
                                                                                                                                                                                                                                                                                                                                                                                                               WO200253774-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Homo
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9; Conservative
                                                                                                                                                                                HENKEL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CAGCAGCCA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (first entry)
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                                                                                                                   Conradt M,
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                                                                                                                       Hofmann
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The invention relates to in vitro identification (M1) of genes expressed in the skin of humans or animals by subjecting a mixture of genetically encoded factors from skin, to serial analysis of gene expression (SACE) so as to identify skin-expressed genes and quantify their expression. (M1) is useful for identifying genes involved in skin homeostasis; to determine skin homeostasis and to test agent (A) that maintains or promotes skin homeostasis or that can be used for treating skin.

Disclosure; Page 88; 1345pp; German.

In vitro identification of a homeostasis and identifying

of skin-expressed genes, useful for ying cosmetic or pharmaceutical agent

agents

determining

skin

cancer.

03-JAN-2001; 2001DE-01000127

HENKEL

Petersohn

'n

Conradt M,

Hofmann

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2002-590638/63

disorders, specifically neurodermatitis; sunburn;

scleroderma;

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ARESULT 69
ABV64486/c
ID ABV644
AC A
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human; skin; dermatological; vulnerary; antipsoriatic; antiseborrhaeic; immunosuppressive; antiinflammatory; cytostatic; SAGE; neurodermatitis;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           11-JUL-2002
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RESULT 70
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Matches 9
                                                        Query Match
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                                                                                                                                         The invention relates to in vitro identification (M1) of genes expressed in the skin of humans or animals by subjecting a mixture of genetically encoded factors from skin, to serial analysis of gene expression (SAGE) so as to identify skin-expressed genes and quantify their expression. (M1) is useful for identifying genes involved in skin homeostasis; to determine skin homeostasis and to test agent (A) that maintains or promotes skin homeostasis or that can be used for treating skin disorders, specifically neurodermatitis; sunburn; psoriasis; scleroderma; ichthyosis; atopic dermatitis; acne; seborrhea; lupus erythematosus; rosacea; melanoma; basal cell carcinoma; and carcinoma or sarcoma of the skin. The present sequence is that of a human expressed sequence tag
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9; Conserv
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TTCAGCAGC 1506
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RESULT 71
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XX ABV67707;
XX 21-OCT-200
XX Human skin
XX Human; skin
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XX Homo sapie
XX Homo sapie
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XX 11-JUL-200
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RESULT 72
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                                                      25-MAR-2003
26-JUL-1994
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                                                                                                                                                                                                         AAQ57385 standard; mRNA;
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07-DEC-1992;
07-DEC-1992;
07-DEC-1992;
19-JAN-1993;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    This is a ACE mRNA target sequence (nucleotide no. 2076) of an enzymatic RNA molecule (ribozyme) which cleaves mRNA associated with the development or maintenance of a cardiovascular condition. The concn. of the ribozyme necessary to effect a therapeutic treatment is lower than that of an antisense oligonucleotide and the specificity of action is higher. (Updated on 25-MAR-2003 to correct PN field.)
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                                                                                                                                                                                                                                                                   SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; peptide nucleic acid; cytosine methylation; cardiovascular; primer; central nervous system; gastrointestinal; respiratory; immune; metak
                                                                                                                                                                                                                                                                                                                                                                           Oligonucleotide primer SEQ ID NO 355765 for detecting
07-APR-2000; 2000DE-01019173
                                                  06-APR-2001; 2001WO-IB000713
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                            range of diseases including immune system, gastrointestinal, respiratory central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC99989, ABF00010-ABH99989 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
                                                                                                                                                                                                                                             This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a
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RESULT 77 ABI77576

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RESULT 76
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                                                                                                                                                                                                                                                                                                                                                                                           This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Set of oligonucleotides, useful for diagnosis and cell typing, designed to detect single-nucleotide polymorphisms and cytosine methylation status.
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                                                                                                                                                                         Sequence
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                                                                                  Score 9; DB 1; Pred. No. 51; 0; Mismatches
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                                                                                                           DB 1
5. 51;
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cardiovascular;
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RESULT 78
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   Synthetic
                                                Eucalyptus; urophylla; grandis; wood volume marker; RAPD genetic marker; random amplified polymorphic DNA analysis; woody perennial plant; family selection; pedigree; mapping; primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 12 BP; 5 A; 4 C; 1 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       auld (FNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are the feet of the control of the co
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Claim 1; SEQ ID NO 377549; 29pp + Sequence Listing; German.
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                                                                                                                                                                            Eucalyptus
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                                                                                                                                                                                                                                                                                              AAQ99836;
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RESULT 79
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XX Yeast
XX Sacch
OS Synth
XX WO983
XX Synth
XX WO983
XX WO981
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Best Local Similarity
Watches 9; Conserve
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Yeast transcriptome - useful for screening antifungal agents, and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 10 BP; 3 A; 4 C; 3 G; 0 T; 0 U; 0 Other;
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                                                                                 WPI; 1998-427943/36.
                                                                                                                                                                                                                                                                                                                                           30-JUL-1998
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Saccharomyces
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Yeast tag
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AAV50097 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   amplifying wood volume markers from Eucalyptus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            fragments from the genomic DNA of E.urophylla, E.grandis and F1 probtained by crossing the two species. Subsequent mapping analysis that the half-sib primers in AAQ99834-Q99840 are all useful for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        RAPD analysis was used to determine whether certain quantitative traits were heritable oligogenic traits in Eucalyptus trees. Sets of commercially available random 10-mer primers were used to amplify
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Example 6; Page 58; 103pp; English.
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                                                                                                                                    Velculescu
                                                                                                                                                                                   (UYJO ) UNIV JOHNS HOPKINS SCHOOL MEDICINE.
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                                                                                                                               Vogelstein
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Pred. No. 56;
                                                                                                                                  Kinzler KW
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  modulating eukaryotic cell, for for identifying genes in cell cycle
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RESULT 80
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Yeast transcriptome is encoded by a DNA molecule comprising a yeast gene involved in cell cycle progression selected from the group of nonannotated ORF (NORF) genes. SAGE (serial analysis gene expression) tags for highly expressed genes and NORF genes are given in AAV50051 to AAV50345. The present invention describes: (1) a method of using yeast genes to modulate the cell cycle which comprises administering to a cell genes to modulate the cell cycle which comprises about the cell cycle which comprises and not be comprised to method of using to a cell genes to modulate the cell cycle which comprises and not be comprised to the cell cycle which comprises and not be comprised to the cell cycle which comprises are given to describe the cell cycle which comprises are given to describe the cell cycle which comprises are given to describe the cell cycle which comprises are given to describe the cell cycle which comprises are given to describe the cell cycle which comprises are given to describe the cell cycle which comprises are given in the cycle which cycle which cycle which cycle which cycle is the cycle which cycle whic
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Kuhlman EG;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    or as an array of probes on a solid support
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90.0%;
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                                                                                               English.
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Pred. No. 56;
                                                                                                                                                                                                                                                                                                          Amerson
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                                                                                                                                                                             rust disease
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The specification describes a method of identifying a genetic marker associated with a genetic locus conferring resistance to fusiform rust disease in a family of trees of the genus Pinus. The method comprises

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RESULT 81
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                                                                                                                 This is a commercial oligonucleotide primer for the PCR-based RAPD analysis of fungal isolates of Dydimella bryiniae and Phoma species. The new method may be used to distinguish D. bryoniae from non-pathogenic Phoma species fungal infections in Cucurbits (i.e.cucumbers, pumpkins, watermelons, gourds, cantaloupes, squashes and related plants). The new method of detection of D. bryoniae and Phoma species infections allows rapid diagnosis even before symptoms are visible as compared to prior art methods which involved growing pure cultures of the pathogens from the infected plants and identifying them under the light microscope. The method leads to the early treatment of the infected plants with
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    16-MAR-1998;
22-FEB-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        recombinant techniques.
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                                                                                        resulting
                                                                                                                                                                                                                                                                                                                                                                                                                                                            Page 8; 1.9pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   assay; test; fungal pathogen;
; RAPD analysis; isolate; food
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                                                                                              ij.
                                                                                              an increased chance of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Dean RA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                            English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ВÞ
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Score 8.4;
Pred. No. 5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          56;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      crop protection; cucurbit;
crop; ss.
                                                                                              saving
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Length 10
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                                                                                                 the infected food
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RESULT 82
AAZ79223
AID AAZ7923
AID AAZ79223
AID AAZ79223
AID AAZ79223
AIC AAZ79
AIC AA
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Best Local Similarity
Matches 9; Conserv
                                                                                                                                                                                                                                                                                                                 19-JUN-1998;
19-JUN-1998;
19-JUN-1998;
19-JUN-1998;
19-JUN-1998;
19-JUN-1998;
19-JUN-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       19-JUN-1998;
19-JUN-1998;
19-JUN-1998;
19-JUN-1998;
19-JUN-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SAGE
APC;
Isolated polynucleotides differentially expressed in cells, useful in gene vaccines against cancer.
                                                                WPI; 2000-106077/09.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Human
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAZ79223 standard; DNA; 10 BP.
                                                                                                                                               (GENZ )
(ROBE/)
(SHAN/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          19-JUN-1998
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   cytotoxic T-lymphocyte; tumour antigen; immunotherapy; anticancer; ss
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     10-APR-2000 (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SAGE tag; serial analysis of gene expression; antigen-presenting cell; APC; monocyte-derived dendritic cell; differential gene expression; immunostimulatory cofactor; costimulatory factor; CTL;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            19-JUN-1998
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1488 GAAGCCAGAC 1497
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           dendritic cell SAGE tag,
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                                                                                                                                               GENZYME CORI
ROBERTS B L.
SHANKARA S.
                                                                                                        BL,
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                                                                                                        Shankara
                                                                                                                                                                                                                                                                                                                 98US-0090036P

98US-0090040P

98US-0090041P

98US-0090042P

98US-0090044P

98US-0090044P

98US-0090044P

98US-0090047P

98US-0090076P

98US-0090076P

98US-0090077P
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98US-0090079P.
98US-0090080P.
98US-0111715P.
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98US-0089997P.
98US-0089999P.
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98US-0089993P
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98US-0089991P
                                                                                                                                                                   CORP.
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Pred. No. 56;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SEQ ID NO:1651.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           DB 1;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Indels
                   antigen-presenting
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0;
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XX Sequences AAZ77573-Z79709 represent SAGE (serial analysis of gene expression) tags used to identify mRNA transcripts encoding commonstimulatory cofactor proteins which are preferentially or differentially expressed in monocyte-derived dendritic cells compared with monocytes. Some of the transcripts correspond to known genes or ESTs (expressed sequence tags) which were previously unknown to be preferentially or differentially expressed in dendritic cells, while clear transcripts correspond to novel genes. Antigen-presenting cell (APC)-associated costimulatory factors play an important role in the cytotoxic immune response, particularly against tumour cells. Tumour antigen presentation via the MRC (major histocompatibility complex) and subsequent recognition by T-cell receptors is alone cefficient to activate a robust cytotoxic immune response that can lyse insufficient of cytotoxic T-lymphocytes (CTs). Nucleic acid cefficient activation of cytotoxic T-lymphocytes (CTs). Nucleic acid cadd complex) and subsequent recognition by T-cell receptors is alone cefficient activation of cytotoxic T-lymphocytes (CTs). Nucleic acid complex) may be used in vaccines to induce an immune response, particularly cagainst a tumour antigen; to modulate the genotype of an APC; to screen in the modulate expression of differentially expressed genes in the modulate expression of differentially expressed genes for the expression of their encoded proteins, can be used to identify cells as belonging to the monocyte lineage. Cells containing these genes can be used to identify expressed genes, or of their encoded proteins, can be used to identify cells as belonging to the monocyte lineage. Cells containing these genes can be used to identify expressed genes in the monocyte lineage. Cells containing these genes cells and vectors containing these genes cells and vectors containing the production of a containing the production of a containing the production of a containing the production of cells containing the production of cells containin
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RESULT 83
AAZ82715
ID AAZ82
XX
AC AAZ8.
DT 07-A
XX
DE Meta
XX
KW Huma
KW Honn
KW ant
XX
PN WOS
XX
PD 23
PR 15
PR 15
PR 15
PR 11 밁 Ş Query Match
Best Local Similarity
Matches 9; Conserv 19-JUN-1998; 19-JUN-1998; 19-JUN-1998; 19-JUN-1998; Human; metastatic breast tumour tissue; breast cancer; tag; primer; non-metastatic breast tumour tissue; gene therapy; anticancer; antimetastatic; vaccine; diagnosis; ss. AAZ82715; AAZ82715 standard; DNA; 10 Metastatic breast tumour cell upregulated transcript tag #1949. 07-APR-2000 23-DEC-1999 WO9965928-A2 18-JUN-1999; 1488 GAAGCCAGAC 1497 sapiens Conservative (first entry) 98US-0089853P. 98US-0089997P. 98US-0090039P. 99WO-US013647 28.0%; <u>,,</u> Score 8.4; D Pred. No. 56; Mismatches BB ۲, ۲. Length 10; Indels 0 Gaps

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RESULT 84
AAZ81205/c
ID AAZ812
XX AAZ812
XX AAZ812
XX O7-APR
XX Metast
XX Human;
KW non-me
KW antime
KW antime
XX Homo s
XX Ho
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Best Local S
Matches 9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 e.g. therapeutic genes (also ribozymes or antisense sequences), particularly an antigen-encoding sequence for use in gene or cell-based vaccines. Polypeptides encoded by the transcripts are also useful in vaccines; for diagnosing breast cancer and for raising specific antibodies (Ab). Ab are used to detect the polypeptides or as therapeutic agents. Host cells that produce the polypeptides can be used to expand and isolate populations of educated, antigen-specific immune effecter cells, e.g. cytotoxic T lymphocytes, and these used for adoptive
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Compounds that modulate expression of the transcripts are potentially useful for treatment of (metastatic) breast cancer, while promoters from the transcripts are used to direct expression, in selected cell types, of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            to AAZ86677 represent tags corresponding to distinct transcripts that are preferentially transcribed in the primary or non-metastatic breast tumour tissue (i.e. are downregulated in metastatic breast tumour cells). These transcripts can be used for diagnosis, prognosis, monitoring and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAZ80767 to AAZ83941 represent tags corresponding to distinct transcripts that are preferentially transcribed in the metastatic breast tumour tissue (i.e. are upregulated in metastatic breast tumour cells). AAZ83942
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Claim 1; Page 111; 219pp; English.
                                                                                                                                                                                                                             WO9965928-A2
                                                                                                                                                                                                                                                                                     Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                Human; metastatic breast tumour tissue; breast cancer; tag; primer; non-metastatic breast tumour tissue; gene therapy; anticancer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 10 BP; 4 A; 4 C; 1 G; 1 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 treatment of breast cancer, particularly where metastatic. Diagnosis is by standard immunoassays or hybridisation/amplification reactions.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Isolated polynucleotides differentially expressed between metastatic and non-metastatic breast cancer cells, useful for diagnosis, prevention and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WPI; 2000-106079/09
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      19-JUN-1998;
19-JUN-1998;
                                                                                                 18-JUN-1999;
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                                                                                                                                                               23-DEC-1999.
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(ROBE/) ROBERTS B L.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (NAHE)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1499 TCAGCAGCCA 1508
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TCAACAGCCA 10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  breast
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Shankara
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (first entry)
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      98US-0089853P.
98US-0089997P.
                                                                                                 99WO-US013647.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        tumour cell upregulated transcript tag #439.
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90.0%;
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Pred. No. 56;
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ARESULT 85
AAZ82619
ID AAZ82
XX AAZ82
XX AAZ82
XX O7-AF
XX Humar
XX Humar
XW Humar
XW Homo
XX Homo

antimetastatic;

non-metastatic breast tumour tissue; gene

vaccine; diagnosis;

Human; metastatic breast tumour tissue; breast cancer; tag;

therapy; anticancer;

18-JUN-1999; 23-DEC-1999

99WO-US013647.

WO9965928-A2

Metastatic breast tumour cell upregulated transcript tag #1853.

07-APR-2000

(first entry)

AAZ82619 standard; DNA; 10

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1486 AAGAAGCCAG 1495

AAGAAGGCAG

Query Match Best Local Matches

Similarity 9; Conserv

28.0%;

Score 8.4; I Pred. No. 56; Mismatches

DB 1;

Length 10;

1.

0

Gaps

0

Conservative

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CC that are preferentially transcribed in the metastatic breast tumour colls, that are preferentially transcribed in the metastatic breast tumour cells). AAZ83942 cc tissue (i.e. are upregulated in metastatic breast tumour cells) that are preferentially transcribed in the primary or non-metastatic breast tumour cells). These cc transcripts can be used for diagnosis, prognosis, monitoring and compounds that modulate expression of the transcripts are potentially where metastatic. Diagnosis is by standard immunoassays or hybridisation/amplification reactions. Cc compounds that modulate expression of the transcripts are potentially. Cc useful for treatment of (metastatic) breast cancer, while promoters from cc the transcripts are used to direct expression, in selected cell types, of ce.g. therapeutic genes (also ribozymes or antisense sequences), compounds that modulate expression of the transcripts are used to direct expression, in selected cell types, of ce.g. therapeutic genes (also ribozymes or antisense sequences), of cc e.g. therapeutic genes (also ribozymes or antisense sequences), of compounds that genes (also ribozymes are also useful in cc vaccines; for diagnosing breast cancer and for raising specific capents. Host cells that produce the polypoptides can be used to expand cc and isolate populations of educated, antigen-specific immune effecter immuncherapy.
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19-JUN-1998;
19-JUN-1998;
Sequence 10 BP; 0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Claim 1; Page 70; 219pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Isolated polynucleotides differentially expressed between metastatic and non-metastatic breast cancer cells, useful for diagnosis, prevention and treatment of cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WPI; 2000-106079/09
                                        immunotherapy
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) ROBERTS B L.
) SHANKARA S.
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98US-0090040P.
98US-0090041P.
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  A;
4 C; 1
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G; 5 T; 0 U; 0 Other;
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19-JUN-1998;

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AAZ84503
ID AAZE
XX
AC AAZE
XX
AC AAZE
DT 07-J
XX
Metc
XX
Metc
XX
Mon
KW Humm
KW non
KW anct
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Hom

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19-JUN-1998;
19-JUN-1998;
                                   23-DEC-1999
                                                                                                                                                                                                                                                Human; metastatic breast tumour tissue; breast cancer; tag; primer; non-metastatic breast tumour tissue; gene therapy; anticancer;
                                                                                                                                                                                                                                                                                                                                      Metastatic breast tumour cell downregulated transcript tag
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAZ84503 standard;
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                                                                                                WO9965928-A2
                                                                                                                                                                                                                      antimetastatic;
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(ROBE/)
(SHAN/)
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ROBERTS B L.
SHANKARA S.
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9; Conserv
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98US-0089997P.
98US-0090039P.
98US-0090040P.
98US-0090041P.
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Pred. No. 56;
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07-APR-2000

(first entry)

AAZ84255;

Human; metastatic breast tumour tissue; breast cancer; tag; primer; Metastatic breast tumour cell downregulated transcript tag #3489.

tumour tissue; gene therapy; anticancer;

Homo sapiens

non-metastatic

breast

WO9965928-A2

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AAZ84255
IID AAZ8
XX
AC AAZ8
AC AAZ8
AC O7-A
DT O7-A
XX
DB Meta
XX
Huma
KW Huma
KW non-
XW anti
XX
OS Homc
XX
PN WO99
XX
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Best Local S
Matches
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19-JUN-1998;
19-JUN-1998;
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(ROBE/)
(SHAN/)
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) ROBERTS B L.
) SHANKARA S.
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9; Conserv
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98US-0089997P.
98US-0090039P.
98US-0090040P.
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Pred. No.
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RESULT 88
AAZ81942/c
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Best Local Similarity
Matches 9; Conserv
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19-JUN-1998;
19-JUN-1998;
                                                            Human; metastatic breast tumour tissue; breast cancer; tag; primer; non-metastatic breast tumour tissue; gene therapy; anticancer:
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(ROBE/)
                                                                                                              Metastatic
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                                              antimetastatic;
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ROBERTS B L.
SHANKARA S.
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                                                                                                            breast
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98US-0089997P.
98US-0090039P.
98US-0090040P.
98US-0090041P.
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                                             vaccine;
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                                                                                                           tumour cell upregulated transcript
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                                             tumour tissue; gene therapy; anticancer;
ne; diagnosis; ss.
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RESULT 89 AAA11232

AAA11232 standard; DNA; 10

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10

CCAGGAAGCC

Matches Query Match Best Local :

l Similarity 9; Conserv

Conservative

0; Mismatches

28.0%;

Pred. Score

8.4; NO. 56 DB 1;

Length 10; Indels

0

Gaps

0

1484 CCAAGAAGCC 1493

PCR primer;

25-OCT-2000

(first entry)

RAPD primer U-12 for detecting Leishmania species.

Identification; Leishmania; isoenzyme; amplicon; promastigote; diagnosis;

amplification; epidemiology; vaccine; ss.

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AAZ80767 to AAZ83941 represent tags corresponding to distinct transcripts that are preferentially transcribed in the metastatic breast tumour tissue (i.e. are upregulated in metastatic breast tumour cella). AAZ83942 to AAZ86677 represent tags corresponding to distinct transcripts that are preferentially transcribed in the primary or non-metastatic breast tumour tissue (i.e. are downregulated in metastatic breast tumour cells). These transcripts can be used for diagnosis, prognosis, monitoring and treatment of breast cancer, particularly where metastatic. Diagnosis is by standard immunoassays or hybridisation/amplification reactions. Compounds that modulate expression of the transcripts are potentially useful for treatment of (metastatic) breast cancer, while promoters from the transcripts are used to direct expression, in selected cell types, of
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19-JUN-1998;
19-JUN-1998;
19-JUN-1998;
                                                     e.g. therapeutic genes (also ribozymes or antisense sequences), particularly an antigen-encoding sequence for use in gene or cell-based vaccines. Polypeptides encoded by the transcripts are also useful in vaccines; for diagnosing breast cancer and for raising specific antibodies (Ab). Ab are used to detect the polypeptides or as therapeuti agents. Host cells that produce the polypeptides can be used to expand and isolate populations of educated, antigen-specific immune effecter cells, e.g. cytotoxic T lymphocytes, and these used for adoptive
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Sequence 10 BP; 0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Isolated polymucleotides differentially expressed between metastatic non-metastatic breast cancer cells, useful for diagnosis, prevention
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                                    immunotherapy
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (GENZ ) GENZYME COR
(ROBE/) ROBERTS B L
(SHAN/) SHANKARA S.
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98US-0089997P.
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98US-0090040P.
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G; 3 T; 0 U; 0 Other;
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RESULT 90
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            identifying isoenzymes and/or amplicons from, respectively, proteins or DNA extracted from cultured promastigotes derived from an isolate. The method uses a limited set of isoenzymes and/or amplification primers that are discriminatory for characteristics of a broad spectrum of Leishmania species. The isoenzymes used for detection include Glucose phosphate Isomerase (GPI), Mannose Phosphate Isomerase (MPI), Nucleoside Hydrolase substrate deoxyinosine (NHD) and Phosphoglucomutase (PGM). The amplification primers used for PCR identification include primers AAA11232 All232. The method is used to identify specific Leishmania species, either for diagnosis or for selection of appropriate treatments, also in epidemiological studies and for preparation of pharmaceuticals
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Identifying species of Leishmania, useful e.g. for diagnosis selection of treatment, using a limited set of isoenzymes or amplification primers for differentiation.
                                                                                                                                                                                          24-NOV-1999;
                                                                                                                                                                                                                                                21-NOV-2000; 2000WO-US031922
                                                                                                                                                                                                                                                                                                      31-MAY-2001.
                                                                                                                                                                                                                                                                                                                                                                                                                                                              Human; transcriptome; gene expression pattern; cancer; drug cancer diagnosis; cell specific gene expression; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Human colon epithelium specific transcriptome sequence SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      20-SEP-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAH63301;
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                            WPI; 2001-367706/38
                                                                                  Velculescu VE,
                                                                                                                                                                                                                                                                                                                                                           WO200138577-A2
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                                                                                                                                       UNIV JOHNS HOPKINS
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                                                                               Vogelstein B,
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                                                                            The present invention describes a method of identifying the type of cell in a sample, involving determining which of the sequences AAH63161-AAH64724 is expressed by the cell. The transcriptomes described in the invention are cell-type specific, cancer specific or ubiquitously expressed in humans. They can also be used to screen for drugs, reduce cancer specific gene expression, standardise expression and restore the function of a diseased cell or tissue. The present sequence is one of the transcriptomes described in the exemplification of the invention
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    Sequence 10
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cell specific gene expression; ss.
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Pred. No. 56;
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Query Match

28.0%;

Score 8.4;

DB

1;

Length 10;

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RESULT 93
ABA06059/c
ID ABA060
XX
AC ABA060
XX
AC ABA060
XX
DT 10-JAN
XX
DE Human
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AAH64237/c
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Matches
                                                                                                                                                                                                   Matches
                                                                                                                                                                                                                Best
                                                                                                                                                                                                                            Query Match
                                                                                                                                                                                                                                                                                                             The present invention describes a method of identifying the type of cell in a sample, involving determining which of the sequences AAH63161-AAH64724 is expressed by the cell. The transcriptomes described in the invention are cell-type specific, cancer specific or ubiquitously expressed in humans. They can also be used to screen for drugs, reduce
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Human;
                                                                                                                                                                                                                                                    Sequence 10 BP; 3 A; 2 C; 3 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                         cancer specific gene expression, standardise expression and restore the function of a diseased cell or tissue. The present sequence is one of the transcriptomes described in the exemplification of the invention
                                                                                                                                                                                                                                                                                                                                                                                       Claim 13; Page 63; 94pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                           New isolated polynucleotides, useful for identifying specific cell type such as cancer cell, comprises transcriptomes expressed in particular
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WO200138577-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAH64237 standard; cDNA; 10
              Human normal
                                      10-JAN-2002
                                                                                      ABA06059 standard; cDNA; 10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WPI; 2001-367706/38
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Velculescu
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         24-NOV-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                21-NOV-2000; 2000WO-US031922
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         31-MAY-2001.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Human
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              20-SEP-2001
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (UYJO ) UNIV JOHNS HOPKINS.
                                                                                                                                                                                                               Local
                                                                                                                                                                            1490 AGCCAGACTT 1499
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ubiquitously
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                                                                                                                                                                                                   similarity
9; Conserve
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                transcriptome; gene expression pattern; cancer; diagnosis; cell specific gene expression; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Similarity 90.09; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TCAACAGCCA 10
                                                                                                                                                    AGCCAGTCTT
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                                                                                                                                                                                                    Conservative
            hepatocyte expression gene
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (first entry)
                                     (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Vogelstein B,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     expressed transcriptome sequence SEQ ID NO:
                                                                                                                                                                                                                28.0%;
90.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       90.0%;
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                                                                                      ВP
                                                                                                                                                                                                   0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        0;
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0; Mismatches
                                                                                                                                                                                                                          Score 8.4;
                                                                                                                                                                                                                Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Kinzler
                                                                                                                                                                                                    Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ₩,
            CDNA,
                                                                                                                                                                                                                          DB 1; Length 10;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1;
            SEQ ID NO: 36
                                                                                                                                                                                                   1;
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                                                                                                                                                                                                    Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             drug screening;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        0
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ARESULT 94
AAF38816
ID AAF38
AC AAF38
XX Yeast
XX Yeast
KW Yeast
KW nor 1
KW nor 1
KW Serid
KW Serid
KW 11inke
XX Sacch
XX 1020
PN W0200
XX 10-U
PN 10-U
XX 10
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Saccharomyces cerevisiae.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Yeast; Saccharomyces cerevisiae; characterisation; cell cycle; NORF; nor previously assigned open reading frame; nonannotated ORF; SAGE; serial analysis of gene expression; antifungal; tag; identification;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Yeast NORF gene SAGE tag oligonucleotide SEQ ID
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           23-MAR-2001
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 10 BP; 1 A; 3 C; 3 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Claim 1; Page 6; 26pp; Japanese
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                                                                                                                                                                                                                                                             Velculescu V,
                                                                                                                                                                                                                                                                                                                                                                                                                                          16-JUN-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      21-DEC-2000.
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                                                                                                                                                                            2001-061874/07.
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                                                                                                                                                                                                                                                                                                                                                        UNIV JOHNS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    primer;
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                                                                                                                                                                                                                                                             Vogelstein B,
                                                                                                                                                                                                                                                                                                                                                                                                                                          99US-00335032
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      gene expression; hepatopathy;
                                                                                                                                                                                                                                                                                                                                                        HOPKINS
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Pred. No. 5
                                                                                                                                                                                                                                                                  Kinzler K;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Mismatches
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Yeast gene coding sequences comprising NORF genes with serial gene expression (SACE) tags, useful for studying, monitoring affecting phases of the cell cycle.

analysis of

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AAF39726
ID AAF39
XX AAF39
XX AAF39
XX AAF39
XX Yeast
XX Yeast
XX Yeast
XX Inke
XX Ink
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Yeast; Saccharomyces cerevisiae; characterisation; cell cycle; NORF; NOR previously assigned open reading frame; nonannotated ORF; SAGE; serial analysis of gene expression; antifungal; tag; identification; linker; PCR primer; ds.
                                             WPI; 2001-061874/07
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WO200077214-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Saccharomyces cerevisiae
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            23-MAR-2001 (first entry)
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                                                                                                                                   Velculescu V,
                                                                                                                                                                                                                                                                                                                                                                                             14-JUN-2000; 2000WO-US016223
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                                                                                                                                                                                                                     UNIV JOHNS HOPKINS.
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                                                                                                                               Vogelstein B,
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                                                                                                                                   Kinzler K;
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The present invention describes an isolated DNA molecule comprising a coding sequence of a yeast gene selected from a group of 745 NORF (not previously assigned open reading frame; or nonannotated ORP) genes CC comprising a SAGE (serial analysis of gene expression) tag. Also Cdescribed are: (1) a method (M1) of using NORF genes to affect the cell Cycle comprising administering a NORF gene whose expression varies by at CC least 10% between any two phases of the cell cycle selected from log phase, S phase and G2/M; (2) a method (M2) for screening candidate antifungal drugs comprising: (a) contacting a test substance with a yeast cell; and (b) monitoring expression of a NORF gene whose expression of the yeast gene is a candidate antifungal drug; (3) a method (M3) for comprising contacting human plan with a probe which modifies the expression of comprising contacting human DNA with a probe which comprises at least 10 contiguous nucleotides of a NORF gene whose expression in a class of drugs having a characteristic effect on gene expression in a cyeast cell comprising contacting a yeast cell with a candidate drug as a member of a class of drugs having a characteristic effect on gene expression in a cyeast cell of at least 1 NORF gene whose expression in a ffected by the class of drugs. The NORF genes may be used to study, monitor and affect phases of the cell cycle, the differentially cycle and for identification of antifungal drugs. AAF33268 to AAF34064 crepressent SAGE tags used in the exemplification of the present invention.

AAF33262 to AAF33267 represent linkers and PCR primers used in the SAGE method, in the exemplification of the present invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Yeast gene coding sequences comprising NORF genes with serial analysis of gene expression (SAGE) tags, useful for studying, monitoring and affecting phases of the cell cycle.
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밁 Ś Best Local Similarity
Matches 9; Conserv Query Match Sequence 10 BP; 4 A; 2 C; 4 G; 0 T; 0 U; 0 Other; 1487 AGAAGCCAGA 1496 \_ AGGAGCCAGA 10 Conservative 28.0%; 90.0%; 0; Mismatches Score 8.4; I Pred. No. 56; DB 1; Length 10; 0, Gaps

0

RESULT 96 AAF40474 AAF40474 standard; DNA; 10 BP

Yeast NORF 23-MAR-2001 (first entry) AAF40474; gene SAGE tag oligonucleotide SEQ

Yeast; Saccharomyces cerevisiae; characterisation; cell cycle; NORF nor previously assigned open reading frame; nonannotated ORF; SAGE; serial analysis of gene expression; antifungal; tag; identification

IJ

NO:7213

Saccharomyces cerevisiae

WO200077214-A2

21-DEC-2000.

14-JUN-2000; 2000WO-US016223.

16-JUN-1999; 99US-00335032

(UYJO ) UNIV JOHNS HOPKINS

Velculescu V, Vogelstein B, Kinzler

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RESULT 97
AAF40855
ID AAF40
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Cycle comprising administering a NORF gene whose expression varies by at CC least 10% between any two phases of the cell cycle selected from log CC phase, S phase and G2/M; (2) a method (M2) for screening candidate CC antifungal drugs comprising; (a) contacting a test substance with a yeast Cc cell; and (b) monitoring expression of a NORF gene whose expression of CC varies as in M1, where a test substance which modifies the expression of the yeast gene is a candidate antifungal drug; (3) a method (M3) for CC identifying human genes which are involved in cell cycle progression CC contiguous nucleotides of a NORF gene whose expression varies as in M1; CC and (4) a method (M4) for identifying a candidate drug as a member of a CC yeast cell comprising contacting a yeast cell with a candidate drug as a member of a CC yeast cell comprising contacting a yeast cell with a candidate drug and CC wonitoring expression in the yeast cell of at least 1 NORF gene whose expression is affected by the class of drugs. The NORF genes may be used to study, monitor and affect phases of the cell cycle, the differentially CC expressed genes may be used as markers of phases of the cell cycle. The CC represent SAGE tags used in the exemplification of the present invention. CC AAF3362 to AAF3367 represent linkers and PCR primers used in the SAGE method, in the exemplification of the present invention.
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Best Local S
Matches 9
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  The present invention describes an isolated DNA molecule comprising coding sequence of a yeast gene selected from a group of 745 NORF (1 previously assigned open reading frame; or nonannotated ORF) genes
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                                                                                                                                                                    Saccharomyces cerevisiae
                                                                                                                                                                                                                         nor previously assigned open reading frame; nonannotated ORF; serial analysis of gene expression; antifungal; tag; identific
                                                                                                                                                                                                                                                                                                                                                                                AAF40855;
                                                                                                                                                                                                                                                                                                                                                                                                                     AAF40855 standard; DNA; 10
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                 16-JUN-1999;
                                                       14-JUN-2000; 2000WO-US016223.
                                                                                           21-DEC-2000
                                                                                                                                                                                                                                                                 Yeast; Saccharomyces cerevisiae; characterisation; cell cycle; NORF;
                                                                                                                                                                                                                                                                                                       Yeast NORF
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9; Conserva
                                                                                                                                                                                                     PCR primer;
                                                                                                                                                                                                                                                                                                   gene SAGE tag oligonucleotide SEQ ID
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Pred. No. 56;
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CC The present invention describes an isolated DNA molecule comprising a CC coding sequence of a yeast gene selected from a group of 745 NORF (not CC previously assigned open reading frame; or nonannotated ORF) genes CC comprising a SAGE (serial analysis of gene expression) tag. Also CC comprising a SAGE (serial analysis of gene expression) tag. Also CC cycle comprising administering a NORF gene whose expression varies by at CC least 10% between any two phases of the cell cycle selected from log CC phase, S phase and G2/M; (2) a method (M2) for screening candidate CC cell; and (b) monitoring expression of a NORF gene whose expression of CC varies as in M1, where a test substance which modifies the expression CC varies as in M2, where a test substance which modifies the expression of the yeast gene is a candidate antifungal drug; (3) a method (M3) for CC contiguous nucleotides of a NORF gene whose expression of CC contiguous nucleotides of a NORF gene whose expression varies as in M1; CC contiguous nucleotides of a NORF gene whose expression varies as in M1; CC contiguous nucleotides of a NORF gene whose expression varies as in M1; CC contiguous nucleotides of a NORF gene whose expression in a CC yeast cell comprising contacting a yeast cell with a candidate drug as a member of a CC peast cell comprising contacting a yeast cell with a candidate drug and md cc yeast cell of at least 1 NORF gene whose cexpression in a problem of the yeast cell of the cell cycle. The cell cycle and for identify candidate drugs. The NORF gene may be used to study, monitor and affect phases of the cell cycle. The cell cycle and for identify candidate drugs which affect the cell cycle. The cell cycle and for identify candidate drugs which affect the cell cycle. The cell cycle and for identify candidate drugs. AAF33268 to AAF44064 crepresent SAGE tags used in the exemplification of the present invention.

CC candidate drugs which affect the cell cycle of the cell cycle. The cell cycle of the cell cycle of the cell cycle of the cell cyc
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밁 á Query Match Best Local : Matches Local 1492 CCAGACTTCA 1501 1 CCAGAGTTCA Similarity 9; Conserv Conservative 10 28.0%; 0 Score 8.4; DI Pred. No. 56; DB 1; ۲. Length 10; Indels <u>,,</u> 0

Sequence 10 BP; 3 A; 3 C; 2 G; 2 T; 0 U; 0 Other;

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RESULT 98
Yeast NORF gene SAGE tag oligonucleotide SEQ ID NO:1941
                                                23-MAR-2001
                                                                                                                                          AAF35202 standard; DNA; 10
                                              (first entry)
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serial analysis of gene expression; linker; PCR primer; ds. Yeast; Saccharomyces cerevisiae; characterisation; cell cycle; NORF; nor previously assigned open reading frame; nonannotated ORF; SAGE; nor previously assigned open reading frame; nonannotated tag;

Saccharomyces cerevisiae

WO200077214-A2

21-DEC-2000

14-JUN-2000; 2000WO-US016223

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CC coding sequence of a yeast gene selected from a group of 745 NORF (not coding sequence of a yeast gene selected from a group of 745 NORF (not previously assigned open reading frame; or nonamnotated ORF) genes CC comprising a SAGE (serial analysis of gene expression) tag. Also CC cleast 10% between any two phases of the cell cycle expression and the yeast cell in the call cycle comprising administering a NORF gene whose expression varies by at CC least 10% between any two phases of the cell cycle selected from log CC phase, S phase and G2/M; (2) a method (M2) for screening candidate cell in the yeast gene is a candidate antifungal drugs comprising; (a) contacting a test substance with a yeast cell; and (b) monitoring expression of a NORF gene whose expression of the yeast gene is a candidate antifungal drug; (3) a method (M3) for comprising contacting human DNA with a probe which comprises at least 10 contifying having a characteristic effect on gene expression in a class of drugs having a characteristic effect on gene expression in a cyeast cell comprising contacting a yeast cell with a candidate drug and monitoring expression in the yeast cell of at least 1 NORF gene whose expression in a cyeast cell comprising contacting a yeast cell with a candidate drug and contacting a propersion of the cell cycle. The C study, monitor and affect phases of drugs. The NORF genes may be used to study the differentially cycle and for identification of antifungal drugs which affect the cell cycle. The cycle and for identification of antifungal drugs which affect the cell cycle. The cycle and for identification of antifungal drugs which affect the cell cycle. The cycle and for identification of antifungal drugs which affect the cell cycle. The method in the exemplification of the present invention.

CC method in the seast of in the exemplification of the present invention.
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Best Local (
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AAF36577
                                                                                               nor previously assigned open reading frame; nonannotated ORF; SAGE; serial analysis of gene expression; antifungal; tag; identification; linker; PCR primer; ds.
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                                                          Saccharomyces cerevisiae
                                                                                                                                                              Yeast; Saccharomyces cerevisiae; characterisation; cell cycle; NORF;
                                                                                                                                                                                                       Yeast NORF gene SAGE tag oligonucleotide SEQ ID NO:3316
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                                                                                                                                                                                                                                                                                                                                standard;
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90.0%;
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Pred.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    core 8.4; DB red. No. 56; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           DB 1; Length 10,
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RESULT 100
AAF36581/c
ID AAF365
XX AAF365
XX AAF365
XX 23-MAR
XX Yeast

Yeast NORF gene SAGE tag oligonucleotide SEQ

ij

NO:3320

Saccharomyces cerevisiae

nor previously assigned open reading frame; nonannotated serial analysis of gene expression; antifungal; tag; iden

tag; identification;

Yeast; Saccharomyces cerevisiae; characterisation; cell cycle; NORF;

AAF36581 standard; DNA; 10

AAF36581;

23-MAR-2001

(first entry)

밁 S

10

CTTGAGCAGC 1

1497 CTTCAGCAGC 1506

Query Match Best Local S Matches 9

Similarity 9; Conserv

Conservative

<u>.</u>

Mismatches

1;

<u>0</u>

Gaps

0

28.0%; 90.0%; 3 C; 3

Score 8.4; Pred. No. 5

DB 1; Length 10 Other;

Sequence

10

BP; 2 A;

G; 2 T; 0 U; 0

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The present invention describes an isolated DNA molecule comprising a CC coding sequence of a yeast gene selected from a group of 745 NORF (not previously assigned open reading frame; or nonannotated ORP) genes CC comprising a SAGE (serial analysis of gene expression) tag. Also CC certification of gene expression tag. Also CC comprising administering a NORF gene whose expression varies by at CC phase. S phase and G2/M; (2) a method (M1) of using NORF genes to affect the cell cycle scattingal drugs comprising: (a) contacting a test substance which a yeast CC cell; and (b) monitoring expression of a NORF gene whose expression of the yeast gene is a candidate antifungal drug; (3) a method (M3) for comprising contacting human genes which are involved in cell cycle progression of the yeast gene is a candidate antifungal drug; (3) a method (M3) for contiguous nucleotides of a NORF gene whose expression of comprising contacting human DNA with a probe which comprises at least 10 contiguous nucleotides of a NORF gene whose expression varies as in M1; and (b) menthod (M4) for identifying a candidate drug as a member of a class of drugs having a characteristic effect on gene expression in a cyeast cell with a candidate drug and monitoring expression in the yeast cell of at least 1 NORF gene whose expressed genes may be used as markers of phases of the cell cycle. The certification of the cell cycle and for identification of the cell cycle. The corresponding to the cell cycle and for identification of the cell cycle. The corresponding to the cell cycle and corresponding to the cell cycle and corresponding to the cell cycle. The corresponding to the cell cycle of the cell cycle of the cell cycle of the cell cycle. The cell cycle and for identification of the cycle of the cell cycle. The cycle and for identification of the cycle of the cell cycle. The cycle and for identification of the cycle of the cell cycle of the cycle
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Velculescu V, Vogelstein B,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              14-JUN-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Example; Page 118; 419pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       16-JUN-1999;
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of.

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CC described are: (1) a method (M1) of using NORF genes to affect the cell Cycle comprising administering a NORF gene whose expression varies by at CC least 10% between any two phases of the cell cycle selected from log CP phase, S phase and G2/M; (2) a method (M2) for screening candidate CC cell; and (b) monitoring expression of a NORF gene whose expression of the yeast gene is a candidate antifungal drug; (a) a method (M2) for screening candidate CC identifying human genes which are involved in cell cycle progression of CC identifying human genes which are involved in cell cycle progression CC comprising contacting human DNA with a probe which comprises at least 10 CC contiguous nucleotides of a NORF gene whose expression in an CC control of the yeast cell comprising contacting a candidate drug; (3) a method (M3) for identifying a candidate drug as a member of a CC class of drugs having a characteristic effect on gene expression in a CC yeast cell comprising contacting a yeast cell with a candidate drug and CC expressed genes may be used to the class of drugs. The NORF gene whose cc expression is affected by the class of drugs. The NORF gene mose cc expression is affected by the class of the cell cycle, the differentially cycle and for identification of antifungal drugs. AAF33268 to AAF44064 CC represent SAGE tags used in the exemplification of the present invention. CC AAF33267 represent linkers and PCR primers used in the SAGE method, in the exemplification of the present invention.
                                                                                                                                                                                                                                                                RESULT 101
                                                                                                                                                                                                                                                                                                                                                                                                               Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
Yeast; Saccharomyces cerevisiae; characterisation; cell cycle; NORF; nor previously assigned open reading frame; nonannotated ORF; SAGE; serial analysis of gene expression; antifungal; tag; identification;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           The present invention describes an isolated DNA molecule comprising coding sequence of a yeast gene selected from a group of 745 NORF (coding sequence of presence of the selected from a group of GRF) genes previously assigned open reading frame, or nonannotated ORF) genes comprising a SAGE (serial analysis of gene expression) tag. Also
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                                                                                       Yeast NORF gene SAGE tag oligonucleotide SEQ ID NO:5639
                                                                                                                                   23-MAR-2001
                                                                                                                                                                              AAF38900
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 10
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                affecting phases of the cell cycle.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                16-JUN-1999;
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                                                                                                                                                                                                                      AAF38900 standard; DNA; 10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (UYJO ) UNIV JOHNS HOPKINS.
                                                                                                                                                                                                                                                                                                                                                                                                                                    Match 28.0%;
Local Similarity 90.0%;
                                                                                                                                                                                                                                                                                                                                                                       1483 ACCAAGAAGC 1492
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    BP; 0 A; 1 C; 3 G; 6 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                  Conservative
                                                                                                                                 (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                          Score 8.4; DB 1; Length 10;
                                                                                                                                                                                                                                                                                                                                                                                                                                          Pred. No. 56;
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                                                                                                                                                                                                                                                                                                                                                                                                                  Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                  1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                  0,
                                                                                                                                                                                                                                                                                                                                                                                                                  Gaps
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RESULT 102
AAF33314
ID AAF333
XX
AC AAF333
XX
DT 23-MAR
XX
DE Yeast
XX

23-MAR-2001 (first entry)

AAF33314 standard; DNA; 10

ΒP

Yeast putative coding sequence SAGE tag oligonucleotide SEQ ID NO:53

Query Match Best Local Similarity Thes 9; Conserva

Conservative

0;

Mismatches

Indels

0

Gaps

0

90.0%;

Score 8.4; D Pred. No. 56;

DB 1; Length 10;

1486 AAGAAGCCAG 1495

Sequence 10 BP; 4 A; 2 C; 4 G; 0 T; 0 U; 0 Other;

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AAGGAGCCAG 10

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The present invention describes an isolated DNA molecule comprising a coding sequence of a yeast gene selected from a group of 745 NORF (not coding sequence of a yeast gene selected from a group of 745 NORF (not comprising a SAGE (serial analysis of gene expression) tag. Also comprising a SAGE (serial analysis of gene expression) tag. Also cycle comprising administering a NORF gene whose expression varies by at cleast 10% between any two phases of the cell cycle selected from log phase, S phase and G2/M; (2) a method (M2) for screening candidate cell; and (b) monitoring expression of a NORF gene whose expression of the yeast gene is a candidate antifungal drug; (a) a method (M2) for screening candidate comprising contacting a test substance which modifies the expression cell; and (b) muman genes which are involved in cell cycle progression of the yeast gene is a candidate antifungal drug; (3) a method (M3) for contiguous nucleotides of a NORF gene whose expression contiguous nucleotides of a NORF gene whose expression varies as in M1; candidate drug as a member of a class of drugs having a characteristic effect on gene expression in a cyeast cell comprising contacting a yeast cell with a candidate drug and monitoring expression in the yeast cell of at least 1 NORF gene whose expression in a feet phases of the cell cycle. The methods may be used to identify candidate drugs which affect the cell cycle and feet phases of the cell cycle. The methods may be used to identify candidate drugs which affect the cell cycle. The method, in the exemplification of the present invention.

Candidate drug sused in the exemplification of the present invention.

Candidate drugs which affect the cell cycle in the call cycle in the cell cycle in the cell cycle. The method, in the exemplification of the present invention.
Yeast gene coding sequences comprising NORF genes with serial analysis gene expression (SAGE) tags, useful for studying, monitoring and affecting phases of the cell cycle.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       linker; PCR primer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Example; Page 201; 419pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (UYJO ) UNIV JOHNS HOPKINS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     9908-00335032
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of.

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RESULT 103
AAF34250
ID AAF342
XX
AC AAF342
XX
XX
AC AAF342
XX
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                                                                                                                                                                                                                                                                                                                                          The present invention describes an isolated DNA molecule comprising a coding sequence of a yeast gene selected from a group of 745 NORF (not previously assigned open reading frame; or nonamnotated ORF) genes comprising a SAGE (serial analysis of gene expression) tag. Also comprising a SAGE (serial analysis of gene expression) tag. Also correct to the cell cycle comprising administering a NORF gene whose expression varies by at CC least 10% between any two phases of the cell cycle selected from log thase, S phase and G2/M; (2) a method (M2) for screening candidate cell; and (b) monitoring expression of a NORF gene whose expression of the yeast gene is a candidate antifungal drug; (3) a method (M3) for centigious nucleotides of a NORF gene whose expression of comprising contacting human DNA with a probe which comprises at least 10 contigious nucleotides of a NORF gene whose expression in a contigious nucleotides of a NORF gene whose expression in a comprising contacting a candidate drug as a member of a CC class of drugs having a characteristic effect on gene expression in a cyeast cell order to ray as a member of a CC class of drugs having a characteristic effect on gene expression in a cyeast cell order in NORF gene whose expression is affected by the class of drugs. The NORF genes may be used to study, monitor and affect phases of the cell cycle, the differentially cycle and for identify candidate drugs which affect the cell cycle and for identification of the cell cycle. The method, M1, and the exemplification of the present invention.

CC chars and for identification of the present invention.

AAF33262 to AAF33267 represent linkers and PCR primers used in the SAGE method, in the exemplification of the present invention.
                                                                                                                                                                                                                                                                              Query Match
Best Local Similarity
  23-MAR-2001
                                         AAF34250;
                                                                                                                                                                                                                                                                                                                                        Sequence 10 BP; 4 A; 3 C; 3 G; 0 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Yeast gene coding sequences comprising NORF genes with serial analysis gene expression (SAGE) tags, useful for studying, monitoring and affecting phases of the cell cycle.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Velculescu V,
                                                                               AAF34250 standard; DNA; 10
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WPI; 2001-061874/07.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          16-JUN-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          linker; PCR primer;
                                                                                                                                                                                                                      1480
                                                                                                                                                                                μ.
                                                                                                                                                                                                                      ACGACCAAGA 1489
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                                                                                                                                                                                                                                                             Conservative
  (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Vogelstein B,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ds.
                                                                                                                                                                                                                                                                              28.0%;
90.0%;
                                                                               BP.
                                                                                                                                                                                                                                                         ٥,
                                                                                                                                                                                                                                                                              Score 8.4; D
Pred. No. 56;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Kinzler K;
                                                                                                                                                                                                                                                             Mismatches
                                                                                                                                                                                                                                                                                                 DB 1; Length 10
                                                                                                                                                                                                                                                         1;
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                                                                                                                                                                                                                                                         Gaps
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Yeast gene coding sequences comprising NORF genes with serial analysis of gene expression (SAGE) tags, useful for studying, monitoring and affecting phases of the cell cycle.
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                                                                                                                                                                                                                                                                                                                                               16-JUN-1999;
                                                                                                                                                                                                                                                                                                                                                                                                              Saccharomyces cerevisiae
                                                                                                                                                                                                                                                                                                                                                                                                                             Yeast; Saccharomyces cerevisiae; characterisation; cell cycle; NORF; nor previously assigned open reading frame; nonamotated ORF; SAGE; serial analysis of gene expression; antifungal; tag; identification; linker; PCR primer; ds.
                                                                                                                                                                                                                                                                                                               Velculescu V,
                                                                                                                                                                                                                                                                                                                               (UYJO ) UNIV JOHNS HOPKINS
                                                                                                                                                                                                                                                                                                                                                               14-JUN-2000; 2000WO-US016223.
                                                                                                                                                                                                                                                                                                                                                                                              WO200077214-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Yeast NORF gene SAGE tag oligonucleotide
                                                                                                                                                                                                                                                                                                               Vogelstein B,
                                                                                                                                                                                                                                                                                                                                               9908-00335032
                                                                                                                                                                                                                                                                                                               Kinzler K;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SEQ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ID NO:989
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The present invention describes an isolated DNA molecule comprising a coding sequence of a yeast gene selected from a group of 745 NORF (not previously assigned open reading frame; or nonannotated ORF) genes comprising a SAGE (serial analysis of gene expression) tag. Also described are: (1) a method (M1) of using NORF genes to affect the cell cycle comprising administering a NORF gene whose expression varies by at least 10% between any two phases of the cell cycle selected from log phase, S phase and G2/M; (2) a method (M2) for screening candidate antifungal drugs comprising: (a) contacting a test substance which a yeast cell; and (b) monitoring expression of a NORF gene whose expression of twaries as in M1, where a test substance which modifies the expression of the yeast gene is a candidate antifungal drug; (3) a method (M3) for identifying human genes which are involved in cell cycle progression contiguous nucleotides of a NORF gene whose expression varies as in M1; and (4) a method (M4) for identifying a candidate drug as a member of a class of drugs having a characteristic affect for case expression in a class of drugs having a characteristic effect on gene expression in a yeast cell comprising contecting a yeast cell with a candidate drug and monitoring expression in the yeast cell of at least 1 NORF gene whose expression is affected by the class of drugs. The NORF genes may be used to study, monitor and affect phases of the cell cycle, the differentially expressed genes may be used as markers of phases of the cell cycle. The methods may be used to identify candidate drugs which affect the cell cycle and for identification of antifungal drugs. ARF33268 to AAF44064 represent SAGE tags used in the exemplification of the present invention. AAF33262 to AAF33267 represent linkers and PCR primers used in the SAGE method, in the exemplification of the present invention

Example; Page 35; 419pp; English.

Sequence 10 BP; 4 A; 3 C; 3 G; 0 T; 0 U; 0 Other;

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Matches
1480 ACGACCAAGA 1489
               Similarity
9; Conserv
                Conservative
                       28.0%;
90.0%;
                0
                              Score 8.4;
                        Pred. No.
                Mismatches
                                 DB 1;
                <u>,</u>
                               Length 10;
                Indels
                0
               Gaps
                0
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RESULT 104 AAF35602/c ID AAF356 XX AAF35602 standard; DNA; 10

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The present invention describes an isolated DNA molecule comprising a coding sequence of a yeast gene selected from a group of 745 NORF (not coding sequence of a yeast gene selected from a group of 745 NORF (not coding sequence of a yeast gene expression) tag. Also comprising a SAGE (serial analysis of gene expression) tag. Also code comprising administering a NORF gene whose expression varies by at collect 10% between any two phases of the cell cycle selected from log cycle comprising administering a NORF gene whose expression varies by at cell; and (b) monitoring expression of a NORF gene whose expression of cell; and (b) monitoring expression of a NORF gene whose expression of the yeast gene is a candidate antifungal drug; (3) a method (M3) for contiguous nucleotides of a NORF gene which expression of comprising contacting human DNA with a probe which comprises at least 10 contiguous nucleotides of a NORF gene whose expression in a yeast cell comprising contacting a characteristic effect on gene expression in a class of drugs having a characteristic effect on gene expression in a yeast cell comprising contacting a yeast cell with a candidate drug and monitoring expression in the yeast cell of at least 1 NORF gene whose expression in a yeast cell comprising contacting a yeast cell with a candidate drug and contacting expression in a present local perfected by the class of drugs. The NORF gene whose expression in a yeast cell with a candidate drug and contacting a present cell with a candidate drug and contacting a candidate of the cell cycle, the differentially expressed genes may be used as markers of phases of the cell cycle. The cell cycle and for identification of antifungal drugs. AAF33268 to AAF44064 represent SAGE tags used in the exemplification of the present invention.

Code methods may be used in the exemplification of the present invention. The method in the exemplification of the present invention.
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                                                                               Query Match
Best Local S
Matches 9
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Yeast gene coding sequences comprising NORF genes with serial analysis gene expression (SAGE) tags, useful for studying, monitoring and
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                                                                                                                                                              Sequence 10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Example; Page 83; 419pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           affecting phases of the cell cycle.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WPI; 2001-061874/07
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    16-JUN-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Yeast NORF gene SAGE tag oligonucleotide SEQ ID NO:2341.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (UYJO ) UNIV JOHNS HOPKINS.
10
                                                                                 Similarity
9; Conser
                                                                                                                                                                                                      in the exemplification of the present invention
                                        CTTCAGCAGC 1506
                                                                                                                                                                ВP;
                                                                                 Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Vogelstein B,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      99US-00335032
                                                                                                                                                                2 A;
                                                                                                                                                                3 C;
                                                                                                  28.0%;
                                                                                                                                                                3 G;
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                                                                                                                       Score 8.4;
                                                                                                    Pred.
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                                                                                                                                                                2 T;
                                                                                 Mismatches
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                                                                                                                                                                0 U;
                                                                                                                       DB 1; Length 10;
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RESULT 105

밁 5

> 1486 AAGAAGCCAG 1495 l Similarity 9; Conserv

Conservative

90.0%; 28.0%; 4

Score 8.4; D. Pred. No. 56;

DB 1; Length 10

۲,

Indels

0

Gaps

0

AAGAAGCCGG

10

Query Match

Local

Sequence

10

ВP;

4. A.

2 C;

G; 0 T; 0 U; 0 Other;

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Coding sequence of a yeast gene selected from a group of 745 NORF (not coding sequence of a yeast gene selected from a group of 745 NORF (not comprising a ssigned open reading frame, or nonannotated ORF) genes comprising a SAGE (serial analysis of gene expression) tag. Also corporate comprising administering a NORF gene whose expression varies by at Cleast 10% between any two phases of the cell cycle screening candidate call intimual drugs comprising: (a) a method (M2) for screening candidate call intimual drugs comprising: (a) a method (M2) for screening candidate call intimual drugs comprising; (a) a method (M2) for screening candidate call intimual drugs comprising; (b) a method (M2) for screening candidate call intimual drugs comprising; (c) a method (M3) for comprising contacting human gene whose expression of comprising contacting human DNA with a probe which comprises at least 10 comprising contacting human but whose expression varies as in M1; candidate drug and method (M3) for identifying a candidate drug as a member of a class of drugs having a characteristic effect on gene expression in a call cycle propersion in a call cycle propersion in a call comprising contacting a yeast cell with a candidate drug and monitoring expression in the yeast cell of at least 1 NORF gene whose expression in a cycle and for identify candidate drugs. The NORF genes may be used to study, monitor and affect phases of the cell cycle. The cycle and for identification of antifungal drugs. AAF33268 to AAF33267 represent in the exemplification of the present invention.

CAAF33262 to AAF33267 represent linkers and PCR primers used in the SAGE method, in the exemplification of the present invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Yeast gene coding sequences comprising NORF genes with serial analysis gene expression (SAGE) tags, useful for studying, monitoring and affecting phases of the cell cycle.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Yeast; Saccharomyces cerevisiae; characterisation; cell cycle; NORF; nor previously assigned open reading frame; nonannotated ORF; SAGE; serial analysis of gene expression; antifungal; tag; identification; linker; PCR primer; ds.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAF38822 standard; DNA; 10
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  16-JUN-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        14-JUN-2000; 2000WO-US016223
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (UYJO ) UNIV JOHNS HOPKINS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Kinzler K;
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of

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Cycle comprising administering a NORF gene whose expression varies by at CC cycle comprising administering and NORF gene whose expression varies by at CC phase, S phase and G2/M; (2) a method (M2) for screening candidate CC antifungal drugs comprising; (a) contacting a test substance with a yeast Cc cell; and (b) monitoring expression of a NORF gene whose expression of CC varies as in M1, where a test substance which modifies the expression of the yeast gene is a candidate antifungal drug; (3) a method (M3) for CC contiguous nucleotides of a NORF gene whose expression CC contiguous nucleotides of a NORF gene whose expression varies as in M1; CC and (4) a method (M4) for identifying a candidate drug as a member of a CC yeast cell comprising contacting a yeast cell with a candidate drug as a member of a CC varies of drugs having a characteristic effect on gene expression in a CC yeast cell comprising contacting a yeast cell with a candidate drug and CC expression is affected by the class of drugs. The NORF gene may be used CC expressed genes may be used as markers of phases of the cell cycle, the differentially CC expressed genes may be used as markers of phases of the cell cycle. The CC cycle and for identify candidate drugs which affect the cell CC cycle and for identification of antifungal drugs. AAF33268 to AAF34064 CC represent SAGE tags used in the exemplification of the present invention.

CC method, in the exemplification of the present invention.
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AAF41495/c
Best Loc
Matches
                Query Match
Best Local :
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       The present invention describes an isolated DNA molecule comprising a coding sequence of a yeast gene selected from a group of 745 NORF (not previously assigned open reading frame; or nonannotated ORF) genes
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Yeast gene coding sequences comprising NORF genes with serial analysis of gene expression (SAGE) tags, useful for studying, monitoring and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Saccharomyces cerevisiae
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Yeast; Saccharomyces cerevisiae; characterisation; cell cycle; NORF; nor previously assigned open reading frame; nonannotated ORF; SAGE; serial analysis of gene expression; antifungal; tag; identification; linker; PCR primer; ds.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    comprising a SAGE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Example; Page 294; 419pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          affecting phases of the cell cycle.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WPI; 2001-061874/07.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Velculescu V;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Yeast NORF gene SAGE tag oligonucleotide SEQ ID NO:8234
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                  Similarity
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                                                                         ₽P;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Vogelstein B,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AGE (serial analysis of gene expression) tag. Also (1) a method (M1) of using NORF genes to affect the cell
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                28.0%;
90.0%;
                                                                         4 C;
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                                                                         G; 4 T;
                Score 8.4;
Pred. No. 9
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                                                                         0 U; 0 Other;
                    56
                                  DB 1;
                                  Length 10;
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9;

Conservative

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Mismatches

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Gaps

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Query Match Sequence 10

BP; 5 A; 2 C; 2 G; 1 T; 0 U;

28.0%;

Score 8.4;

DB 1; 0 Other;

Length 10;

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ARBSULT 107
AARS14/AXX
XX
AARS14/AXX
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AARS14/AX

                                               The present invention describes an isolated DNA molecule comprising a coding sequence of a yeast gene selected from a group of 745 NORF (not previously assigned open reading frame; or nonannotated ORF) genes comprising a SAGE (serial analysis of gene expression) tag. Also comprising a SAGE (serial analysis of gene expression) tag. Also correct the cell cycle comprising administering a NORF gene whose expression varies by at cell east 10% between any two phases of the cell cycle selected from log contifingal drugs comprising: (a) a method (M2) for screening candidate cell; and (b) monitoring expression of a NORF gene whose expression of cell; and (b) monitoring expression of a NORF gene whose expression of the yeast gene is a candidate antifungal drug; (3) a method (M3) for cell; and (b) muman genes which are involved in cell cycle progression of comprising contacting human DNA with a probe which comprises at least 10 contiguous nucleotides of a NORF gene whose expression varies as in M1; and (4) a method (M4) for identifying a candidate drug as a member of a class of drugs having a characteristic effect on gene expression in a cyeast cell of at least 1 NORF gene whose expression in a cyeast cell of at least 1 NORF gene may be used to study, monitor and affect bases of the cell cycle. The methods may be used to identify candidate drugs. Apr33268 to AAF33267 represent linkers and PCR primers used in the SAGE cethod, in the exemplification of the present invention.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Yeast gene coding sequences comprising NORF genes with serial analysis gene expression (SAGE) tags, useful for studying, monitoring and affecting phases of the cell cycle.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Example; Page 41; 419pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WPI; 2001-061874/07.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Yeast NORF gene SAGE tag oligonucleotide SEQ ID NO:1165
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          9908-00335032
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CC coding sequence of a yeast gene selected from a group of 745 NORF (not Cr previously assigned open reading frame; or nonannotated ORP) genes CC comprising a SAGE (serial analysis of gene expression) tag. Also CC described are: (1) a method (M1) of using NORF genes to affect the cell CC cycle comprising administering a NORF gene whose expression varies by at least 10% between any two phases of the cell cycle selected from log phase, S phase and G2/M; (2) a method (M2) for screening candidate CC antifungal drugs comprising: (a) contacting a test substance with a yeast Cc the yeast gene is a candidate antifungal drug; (3) a method (M3) for comprising contacting human Genes which modifies the expression CC varies as in M1, where a test substance which modifies the expression CC contiguous nucleotides of a NORF gene whose expression of the yeast gene is a candidate antifungal drug; (3) a method (M3) for CC contiguous nucleotides of a NORF gene whose expression varies as in M1; and (4) a method (M4) for identifying a candidate drug as a member of a CC class of drugs having a characteristic effect on gene expression in a CC yeast cell comprising contacting a yeast cell with a candidate drug and CC methods may be used as markers of phases of the cell cycle. The cepressed genes may be used as markers of phases of the cell cycle. The cream tags used in the exemplification of the present invention. CC MAF33262 to AAF33267 represent of the present invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Best Local Similarity 90. Matches 9; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Yeast gene coding sequences comprising NORF genes with serial analysis gene expression (SAGE) tags, useful for studying, monitoring and affecting phases of the cell cycle.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Yeast; Saccharomyces cerevisiae; characterisation; cell cycle; NORF; nor previously assigned open reading frame; nonannotated ORF; SAGE; nor previously assigned open reading frame; nonannotated ORF; SAGE; nor previously assigned open expression; antifungal; tag; identification;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WPI; 2001-061874/07
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAF34234 standard; DNA; 10 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Example; Page 34; 419pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (UYJO ) UNIV JOHNS HOPKINS
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                  exemplification of the present invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Vogelstein B,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        9908-00335032
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Kinzler K;
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XX DE Primez
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KW Growth
KW isolat
XX Homo E
XX WO2001
XX Genot
PI Chew !
XX WPI;
XX Genot
PI indiv:
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                                                                                                                                                                                                                                                                      (SNPs) in the human growth hormone releasing hormone receptor (GHRHR) gene located on chromosome 7p14, and methods for haplotyping and/or genotyping the GHRHR gene. The methods of the invention make use of allele-specific oligonucleotides (ASOs) as probes and primers and/or primer-extension oligonucleotides for detecting the GHRHR gene polymorphisms. The polynucleotides and screened compounds are useful the treatment of diseases associated with GHRHR activity, such as isolated growth hormone deficiency (IGHD) and pituitary adenomas. AAS19648-AAS19673 represent primer-extension oligonucleotides for detecting human GHRHR gene polymorphisms
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                                                                                                                                                                                                                     Sequence 10 BP;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Genotyping human Growth hormone releasing hormone receptor gene of individual for determining haplotype of individual by determining identity of nucleotide pair at specific polymorphic sites for two
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WPI; 2002-066342/09
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9; Conserv
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                                                    GAAGCCAGAC 1497
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    GACTICACCA
GAAGCCTGAC 10
                                                                                                            Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               invention relates to novel single nucleotide polymorphisms
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90.0%;
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                                                                                                                                   Score 8.4; D
Pred. No. 56;
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Pred. No. 5
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                                                                                                            Mismatches
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                                                                                                            Gaps
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ABL52200 standard;

DNA;

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RESULT 111
ARK61399/C
ID ABK613
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AC ABK813
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DT 13-AUG
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           The present invention describes an isolated human period (Drosophila) CC homologue 1, (PER1) polynuclectide (I) comprising a sequence which is a copolymorphic variant for a reference sequence (ABL52077) for the PER1 gene CC or its fragment, or a polymorphic variant of a reference sequence (ABL52078) for a PER1 CDNA or its fragment. The present invention also comparished the present invention also comparished the present invention also compared to the personal protein for use in screening for candidate compared to treat diseases related to PER1 activity. (I) is useful for therapeutic purposes A recombinant non-human organism transformed or therapeutic purposes. A recombinant non-human organism transformed or transfected with (I) can be used for studying expression of the PER1 isogenes in vivo, for in vivo screening and testing of drugs targeted against PER1 protein, and for testing the efficacy of therapeutic agents associated with circadian rhythm regulation. The present sequence represents a preferred oligonucleotide primer for the present sequence represents a preferred oligonucleotide primer for the present sequence represents a preferred oligonucleotide primer for the present sequence represents a preferred oligonucleotide primer for the present sequence represents a preferred oligonucleotide primer for the present sequence represents a preferred oligonucleotide primer for the present sequence represents a preferred oligonucleotide primer for the present sequence represents a preferred oligonucleotide primer for the present sequence represents a preferred oligonucleotide primer for the present sequence represents a preferred oligonucleotide primer for the present sequence represents and the present sequence represents a preferred oligonucleotide primer for the present sequence represents and the present sequence represents and the present sequence represents a preferred oligonucleotide primer for the present sequence represents and the presents and the present sequence represents and the present sequence re
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Best Local S
Matches 9
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                                            SCYA21 gene primer extension oligonucleotide #3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Novel isolated human period Drosophila homolog 1 polynucleotide, useful for therapeutic purposes, for studying the expression and function of the polynucleotide, and for expressing the homolog.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Homo sapiens.
Small inducible cytokine subfamily A (Cys-Cys) member 21; SCYA21;
                                                                                                  13-AUG-2002
                                                                                                                                                   ABK81399
                                                                                                                                                                                                  ABK81399 standard;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Page 16; 162pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      BP; 2 A; 4 C; 3 G; 1 T; 0 U; 0 Other;
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23-MAY-2002

WO200240502-A2

polymorphism;

primer; ss.

sapiens.

Human; lipase; hormone sensitive; LIPE; isogene; obesity; male sterility;

Human LIPE gene polymorphism detection oligonucleotide primer #38

ABK96063;

ABK96063 standard;

DNA;

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24-SEP-2002

(first entry)

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ABK96063/c
ID ABK960
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AC ABK960
AC ABK960
DT 24-SEP
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DE Human
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 10 BP; 0 A; 4 C; 2 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        The invention describes an isolated polynucleotide, which comprises genes and haplotypes of the small inducible cytokine subfamily A (Cys-Cys),
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Claim 16; Page 13; 56pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            09-OCT-2001; 2001WO-US046141.
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drug development; immunomodulator; primer extension; oligonucleotide; ss.
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RESULT 113
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Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              expressing LIPE protein for use in screening for candidate drugs to treat diseases related to LIPE activity. It is also useful in studying the effect of the variation on the biological activity of LIPE as well as on the binding affinity of candidate drugs targeting LIPE for the treatment of obesity and male sterility. The invention is useful for studying the expression of LIPE isogenes in vivo, for in vivo screening and testing of drugs targeted against LIPE protein, and for testing the efficacy of therapeutic agents and compounds for treating obseity and male sterility in a biological system. The present nucleic acid sequence represents one of a collection (ABK96026-ABK96083) of oligonucleotide primers that were
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       isogenes. The invention is useful in screening for drugs targeting LIPE isogenes that are useful for treating obesity and male sterility. The methods of the invention are useful for improving the efficiency and reliability of several steps in the discovery and development of drugs for treating diseases associated with LIPE activity. The polynucleotide is useful in studying the expression and function of LIPE, and in
                                                                                                                                                                                                                                                                                                                                                                                                          Human; homeo box D3; HOXD3; polymorphism; developmental disorder;
haplotype; HT; allele-specific oligonucleotide; ASO; tumour; therapy;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           The present invention relates to a new polynucleotide comprising nucleotide sequence which comprises lipase, hormone sensitive (Li
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Novel genetic variants of Lipase, Hormone-Sensitive isogenes, useful for improving efficiency and reliability in drug development for treating diseases associated with LIPE activity, e.g. obesity and male sterility.
                                                                                                                         24-MAY-2001; 2001WO-US016982.
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      (GENA-)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                box D3 (HOXD3) gene
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Pred. No. 56;
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RESULT 114
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             in a variety of diagnostic and prognostic formats and therapeutic methods. A recombinant non-human organism is useful in studying expression of the EOXID isogenes in vivo. Allele-specific oligonucleotides (ASO) are useful as probes and primers and for assaying a polymorphism in the target region. The present sequence is a primer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SAGE tag; serial analysis of gene expression; human; chronic hepatitis CH; liver tissue; hepatocellular carcinoma; cancer; tumour; HCC;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ABV84823;
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Pred. No. 56;
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Human chronic hepatitis C tissue expression exasperating gene group comprises 100 high-ranking genes.

WPI; 2002-631294/68

19-JAN-2001; 2001JP::00012328

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19-JAN-2001; 2001JP-00012328

KAGAKU GIJUTSU SHINKO JIGYODAN

Claim 55; Page 28; 139pp; Japanese

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RESULT 115
ABK54423/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CC The invention relates to SAGE (serial analysis of gene expression) tags CC representing groups of genes which are differentially expressed in human CC chronic hepatitis C (CH) liver tissue or hepatitis C cinduced (CH) liver tissue or hepatitis C cinduced (CH) liver tissue or hepatitis C cinduced (CH) liver tissue or hepatitis C induced (CH) liver tissue or hepatitis C induced (CH) liver tissue or hepatitis of a sequence of 10 nucleotides (CH) liver tissue of the 5'-CATG-3' sequence motif lying nearest to the CO polya region of cDNAs derived from a variety of genes. These tags serve to uniquely identify each transcript and can thus be used to analyse the CC polya region of cDNAs derived from a variety of genes. These tags serve to uniquely identify each transcript and can thus be used to analyse the CC pattern of gene expression in particular cell types. The invention also CC relates to proteins encoded by the genes expressed in chronic hepatitis C liver tissue or HCC, antibodies against these proteins, and inhibitors of CC the expression of groups of genes that are overexpressed in chronic CC hepatitis C liver tissue or HCC. Groups of genes differentially expressed CC in chronic hepatitis C tissue or HCC may be used for the diagnosis and CC or activity, and antibodies against the gene products may be used in the CC expressed in chronic hepatitis C and/or HCC. Sequences CC ABV84791-ABV84890 are SAGE tags representing 100 genes which are highly cx expressed in chronic hepatitis C liver tissue
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Best Local
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                                                                                                                          expressing ISL
treat diabetes
                                                                                                                                       Novel isolated polynucleotide which is a polymorphic variant of Istranscription factor, LIM/Homeodomain, (islet-1) (ISL1) used for expressing ISL1 protein isoform and for screening drug candidates
                                                                                                                                                                                                                                                                                                                             04-AUG-2000; 2000US-0223535P
                                                                                                                                                                                                                                                                                                                                                                 06-AUG-2001; 2001WO-US024664
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Human; ss; primer; ISL1; islet-1; chromosome 5q; motor neuron defect; diabetes; transcription factor; LIM; homeodomain; antidiabetic; PCR;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human ISL1 gene ASO primer extension sequence #18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ABK54423
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                                                                                                                                                                                                                                                                                            PHARM INC
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Pred. No. !
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The invention relates to an isolated polynucleotide sequence which comprises ISII transcription factor (islet-1, of the LIM/Homeodomain family), isogene and the polymorphic variants of the coding region (cDNA). Also included are a recombinant non-human organism expressing

The present invention relates to a new polypeptide comprising a sequence which is a polymorphic variant of a reference sequence for ADRB3 (beta-3-adrenergic receptor) protein. The reference sequence comprises a sequence

14;

English

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RESULT 116
ABK11491
ID ABK114
AC ABK114
XX ABK114
AC ABK114
XX Oligon
XX Human;
KW antidi
KW single
XX Single
XX W02002
XX HOMO &
PF 23-JUI
XX 21-JUI
XX 16ENA-
XX WPI; :
XX WPI; :
XX WOVel
PT Study:
PT Stud
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                                                                                                                                                                                               Novel genetic variants of beta-3-adrenergic receptor gene useful in studying expression and function of the protein, and for screening drugs to treat diseases e.g. obesity, non-insulin dependent diabetes mellitus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human; beta-3-adrenergic; receptor; ADRB3; primer; anorectic; ss; antidiabettc; gene therapy; morbid obesity; insulin resistance; non-insulin-dependent diabetes mellitus; haplotyping; SNP; single nucleotide polymorphism.
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                                                                                                                                                                                                                                                                                                                                                 WPI; 2002-241571/29.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (GENA-) GENAISSANCE
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                                                                                                                                       Page 15;
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                                                                                                                                       91pp; English
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Pred. No. 56;
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RESULT 117
ABK72629
ID ABK7262
XX ABK726
XX Leukot
XX Leukot
XX Human;
KW Chemoa
KW inflam
KW primer
XX Homo s
XX GENA-
XX Hora;
XX
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The CC polymorphic variants are useful in studying the expression and function CC of ADRB3, in expressing ADRB3 protein for use in screening for candidate CC drugs to treat diseases related to ADRB3 activity, in studying the effect CC of the variation on the biological activity of ADRB3, and the binding CC affinity of candidate drugs targeting ADRB3 for the treatment of CC disorders such as morbid obesity, insulin resistance and an early onset CC on on-insulin-dependent diabetes mellitus. Haplotyping methods are CC useful in validating ADRB3 as a candidate target for treating a specific condition or disease predicted to be associated with ADRB3 activity, or CC in the design of clinical trials of candidate drugs for treating a Specific condition or disease associated with ADRB3 activity. The present CC nucleic acid sequence represents one of a collection of oligonucleotide CC invention to defeor nolvmorphisms in the methods of the CC invention to defeor nolvmorphisms in the human account CC invention to defeor nolvmorphisms in the human account CC invention to defeor nolvmorphisms in the human account CC invention to defeor nolvmorphisms in the human account CC invention to defeor nolvmorphisms in the human account CC invention to defeor nolvmorphisms in the human account CC invention to defeor nolvmorphisms in the human account CC invention to defeor nolvmorphisms in the methods of the CC invention to defeor nolvmorphisms in the human account CC invention to defeor nolvmorphisms in the human account CC invention to defeor nolvmorphisms in the human account control                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
Best Local S
Matches 9
                                 The invention describes an isolated human leukotriene B4 receptor (chemokine receptor-like 1) (LTB4R) polynucleotide (I) comprising a sequence which is a polymorphic variant for a reference sequence for LTB4R gene or its fragment, or a polymorphic variant of a reference sequence for a LTB4R cDNA or its fragment. LTB4R is a potent chemoattractant that is primarily involved in inflammation, immune responses and host defense against infection. (I) is useful in studying the sequence of the seq
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Novel isolated human leukotriene B4 receptor polynucleotide, useful therapeutic purposes, for studying expression and function of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Bieglecki KM,
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                                                                                                                                                                                                                                                                                                                                                                                                                     Claim 17;
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                                                      adenoma), comprising: (a) detecting macrophage inhibitory cytokine (MIC) or remail dipeptidase (RDP) in faeces or blood of a subject and comparing amount of MIC or RDP detected to that in normal subjects, where an elevated amount of MIC or RDP in the subject is an indicator of CC in subject; (b) isolating mRNA sample from faeces of a subject, detecting MIC or RDP mRNA in the mRNA sample, and comparing amount of MIC or RDP mRNA in the subject is an indicator of CC in subject or RDP mRNA in the subject is an indicator of CC in subject; (c) rRDP mRNA in the subject is an indicator of CC in subject; (c) isolating epithelial cells from blood of a subject, isolating an mRNA sample from faeces of a subject or epithelial cells, detecting MIC or RDP mRNA in the mRNA sample, and comparing the amount of MIC or RDP mRNA in the mRNA sample to amounts of MIC or RDP mRNA in the mRNA sample to amounts of MIC or RDP mRNA in the mRNA sample to amounts of MIC or RDP mRNA in the mRNA sample to amounts of MIC or RDP mRNA in the mRNA sample to amounts of MIC or RDP mRNA in the mRNA sample to amounts of MIC or RDP mRNA in the mRNA sample to amounts of MIC or RDP mRNA in the mRNA sample to amounts of MIC or RDP mRNA in the mRNA sample to amounts of MIC or RDP mRNA in the mRNA sample to amounts of MIC or RDP mRNA in the mRNA sample to amounts of MIC or RDP mRNA in the mRNA sample to amounts of MIC or RDP mRNA in the mRNA sample to amounts of MIC or RDP mRNA in the mRNA sample to amounts of MIC or RDP mRNA in the mRNA sample to amounts of MIC or RDP mRNA in the mRNA sample to amounts of MIC or RDP mRNA in the mRNA sample to amounts of MIC or RDP mRNA in the mRNA sample to amounts of MIC or RDP mRNA in the mRNA sample to amounts of MIC or RDP mRNA in the mRNA sample to amounts of MIC or RDP mRNA in the mRNA sample to amounts of MIC or RDP mRNA in the mRNA sample to amounts of MIC or RDP mRNA in the mRNA sample to amounts of MIC or RDP mRNA in the mRNA sample to amounts of MIC or RDP mRNA in the mRNA sample to amount of MIC or RDP mRNA in
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       an elevated of CC in the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Detecting colorectal cancer in a subject, involves detecting macrophage inhibitory cytokine or renal dipeptidase or their mRNA in feces or blood
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                                       the mRNA sample is
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RESULT 119
ABT1441
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XX ABT144
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DT 20-FEE
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XX WO20/
PN WO20/
PN U7-0
XX U7-0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 an RDP substrate, detecting activity of RDP in the blood or faeces by CC detection of increased reaction product or decreased RDP substrate, and CC comparing the amount of activity of RDP in blood or faeces of the subject to that in normal subjects, where an elevated amount of activity of RDP CC in the blood or faeces of the subject is an indicator of CC in the subject; (e) administering to a subject is an intibody or inhibitor is comparing the moiety which is detectable from outside of the subject and CC detecting the moiety which is detectable from outside of the subject and CC area of localisation of the moiety within the subject but outside the cC proximal tubules of the kidney identifies CC; or (f) administering to a subject a substrate for RDp, the substrate being labeled with a CC detectable moiety, isolating faeces or blood from the subject, and CC detectable moiety, where increased product or RDP substrate CC with the detectable moiety, where increased product or decreased CC substrate in the faeces or blood indicates CC in the subject. The methods care useful for detecting colorectal cancer in a subject. The present CC sequence is a DNA tag derived from a human transcript whose expression is clevated in colorectal cancer or colorectal adenoma
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                      The invention comprises a method for the PCR amplification of nucleic acids. The method involves a set of primers, where two of the primers a in solution and at least two other primers are attached to a solid support. The method of the invention can be used for the analysis of a nucleic acid or a mixture of nucleic acids, including: single-stranded DNA molecules, double-stranded DNA molecules, double-stranded DNA molecules and mRNA molecules. The
                                                                                                                                                                                                                                 Amplification of nucleotide sequences from polynucleotides by chain extension of oligonucleotide primers, comprises 2 oligonucleotides in solution, 2 attached to supports and both share complementary sequences.
                                                                                                                                                                                       Disclosure; Fig 17; 60pp; English
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    a random amplified
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    polymorphic DNA (RAPD)
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                                                                                                                                                                                                    The invention comprises a method for the PCR amplification of nucleic acids. The method involves a set of primers, where two of the primers in solution and at least two other primers are attached to a solid support. The method of the invention can be used for the analysis of a nucleic acid or a mixture of nucleic acids, including; single-stranded DNA molecules, double-stranded DNA molecules, double-stranded DNA molecules, and mRNA molecules. The present DNA sequence represents a random amplified polymorphic DNA (RAF PCR primer of the invention
                                                                                                                                                                                Sequence 10
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                                                                                                                                                                                                                                                                                                                                         Amplification of nucleotide sequences from polynucleotides by chain extension of oligonucleotide primers, comprises 2 oligonucleotides solution, 2 attached to supports and both share complementary seques
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Nucleic acid PCR
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nes 9; Conserv
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                                                                                                                                Similarity 9; Conserv
                                                                                                          AAGAAGCCAG
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                                                                                                                                                                                                                                                                                                                  Fig 17; 60pp; English.
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Pred. No. 56;
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                                                                                                                                                                                                                    polymorphic DNA (RAPD)
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                                                                                                                                                                                                                                                                                                                                          sequences.
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AAX14955

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ABQ86554
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           The present sequence represents a polynucleotide that is able to form triple helix with a double stranded sequence. Cytosine bases in the present can be replaced with 5-methylcytosine for increased triplex stability. The present sequence is used in the assay of the invention, where it can be part of the anchor DNA or reporter DNA sequence. The assay comprises adding a sample containing double-stranded DNA test
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Assay of genetic sequences based on triplex formation from double stranded analyte - and hybrid of anchor and reporter sequences, with reporter released if triplex formation occurs, used e.g. to identify
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24-MAR-1999
              Human; skin ageing; skin stress; EST; expressed sequence tag; ss.
                                            Human skin stress/ageing related EST SEQ ID NO 309.
                                                                              10-SEP-2002
                                                                                                             ABQ86554;
                                                                                                                                             ABQ86554 standard; cDNA; 11 BP
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                                                                                                                                                                                                                                                                                                                                                                                      field.)
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                                                                                                                                                                                                                                                                                       Local Similarity 90.
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(first entry)
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RESULT 123
ABQ86306/C
ID ABQ863
XX ABQ863
XX ABQ863
XX Human
XX Human
XX Human;
XX Human;
XX Homo s
XX Y
PN WO2002
XX 11-JUL
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Best Local :
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                                                                                                                                                                                                   03-JAN-2001; 2001DE-01000121
                                                                                                                                                                                                                                                                                                                                                                        WO200253773-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Human; skin ageing; skin stress; EST; expressed sequence tag;
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                                                                                    Petersohn D, Conract M,
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                                                                                    Hofmann K;
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Pred. No. 60;
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Best Local
                Claim 8; Page //; ביבור Claim 8; Page containt for skin ageing and/or skin stress by serial analysis of gene expression between mixtures of transcribed and optionally translated, genetically encoded factors (A) obtained from young and aged skin, to identify that genes that show strong differential expression. (A) comprises protein or mRNAs or their fragments. (M1) is useful for: identifying markers of skin ageing and/or stress; and identifying or determining the effects of the contained of skin ageing. The present
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   skin ageing and/or stress; and identifying or determining the pharmaceutical or cosmetic agents for control of skin ageing. sequence is one of a group of human skin ageing/stress relates
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                                                                                                                                                                                                                                                         The invention relates to in vitro identification (M1) of genes expressed in the skin of humans or animals by subjecting a mixture of genetically encoded factors from skin, to serial analysis of gene expression (SAGE) so as to identify skin-expressed genes and quantify their expression (M1) is useful for identifying genes and quantify their expression. (M1) is useful for identifying genes involved in skin homeostasis; to determine skin homeostasis and to test agent (A) that maintains or promotes skin homeostasis or that can be used for treating skin disorders, specifically neurodermatitis; sunburn; psoriasis; scleroderma; ichthyosis; atopic dermatitis; acne; seborrhe; lupus erythematosus; rosaces; melanoma; basal cell carcinoma; and carcinoma or sarcoma of the skin. The present sequence is that of a human expressed sequence tag (EST) of the invention
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  The invention relates to in vitro identification (M1) of genes expressed in the skin of humans or animals by subjecting a mixture of genetically encoded factors from skin, to serial analysis of gene expression (SACE) so as to identify skin-expressed genes and quantify their expression. (M1) is useful for identifying genes involved in skin homeostasis; to determine skin homeostasis and to test agent (A) that maintains or promotes skin homeostasis or that can be used for treating skin
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Human; skin; dermatological; vulnerary; antipsoriatic; antiseborrhaeic; immunosuppressive; antiinflammatory; cytostatic; SAGE; neurodermatitis; psoriasis; dermatitis; skin cancer; EST; expressed sequence tag; ss.
                                                                                                                                Human skin
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                                                                                                                                                                                                                                                     ABV69450;
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Pred. No. (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                             neurodermatitis;
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03-JAN-2001; 2001DE-01000127

20-DEC-2001; 2001WO-EP015179.

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RESULT 129
ABV66348/c
ID ABV663
XX ABV663
XX Human
XX Human;
KW Human;
KW Humuno
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XX Homo s
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                                                                                                                                                                                                                                                                                                                                                                                               Human; skin; dermatological; vulnerary; antipsoriatic; antiseborrhaeic; immunosuppressive; antiinflammatory; cytostatic; SAGE; neurodermatitis; psoriasis; dermatitis; skin cancer; EST; expressed sequence tag; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Human skin
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                                                            Disclosure; Page 139; 1345pp; German
                                                                                                       In vitro identification of shomeostasis and identifying
                                                                                                                                                       WPI; 2002-590638/63.
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The invention relates to in vitro identification in the skin of humans or animals by subjecting a encoded factors from skin, to serial analysis of

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                                                        The invention relates to in vitro identification (M1) of genes expressed in the skin of humans or animals by subjecting a mixture of genetically encoded factors from skin, to serial analysis of gene expression (SAGE) so as to identify skin-expressed genes and quantify their expression (M1) is useful for identifying genes involved in skin homeostasis; to determine skin homeostasis and to test agent (A) that maintains or promotes skin homeostasis or that can be used for treating skin disorders, specifically neurodermatitis; sunburn; psoriasis; scleroderma; ichthyosis; atopic dermatitis, acne; seborrhea; hupus erythematosus; rosacea; melanoma; basal cell carcinoma; and carcinoma or sarcoma of the skin. The present sequence is that of a human expressed sequence tag (EST) of the invention
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      03-JAN-2001; 2001DE-01000127
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 nomeostasis and identifying cosmetic or pharmaceutical agents against
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ARESULT 131
ARESULT 131
ARV6670
ID ABV667
XX ABV6677
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XX 21-OCT
XX Human;
KW Human;
KW Human;
KW Human;
KW Human;
CX WO2002
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RESULT 132
ABV68413/c
ID ABV684
XX
AC ABV684
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     ABV68413;
                                                         ABV68413 standard; cDNA; 11 BP
                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 11 BP;
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RESULT 133
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XX ABV690
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XX Human
XX Immuno
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                                                                          Human; skin; dermatclogical; vulnerary; antipsoriatic; antiseborrhaeic; immunosuppressive; antiinflammatory; cytostatic; SAGE; neurodermatitis; psoriasis; dermatitis; skin cancer; EST; expressed sequence tag; ss.
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                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 11 BP; 2 A; 2 C; 5
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Pred. No. 60;
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RESULT 134
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ABV683
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AC ABV683
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DT 21-OCT
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Human;
KW Human;
KW immuno
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Homo 8
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HOMO 11-JUL
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skin. The present sequence is
(EST) of the invention
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homeostasis and identifying cosmetic or pharmaceutical agents against
      196;
1345pp;
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The invention relates to in vitro identification (M1) of genes expressed in the skin of humans or animals by subjecting a mixture of genetically encoded factors from skin, to serial analysis of gene expression (SAGE) so as to identify skin-expressed genes and quantify their expression. (M1) is useful for identifying genes involved in skin homeostasis; to determine skin homeostasis and to test agent (A) that maintains or promotes skin homeostasis or that can be used for treating skin discovers Sequence 11 BP; 3 A; 4 C; disorders, specifically neurodermatitis; sunburn; psoriasis; sclerode ichthyosis; atopic dermatitis; acne; seborrhea; lupus erythematosus; rosacea; melanoma; basal cell carcinoma; and carcinoma or sarcoma of skin. The present sequence is that of a human expressed sequence tag (EST) of the invention ω G; 1 T; 0 U; 0 Other; scleroderma; of the

Matches Query Match Best Local ( 1484 N 9 Similarity CCAAGATGCC 11 CCAAGAAGCC Conservative 1493 28.0%; 0 Pred. No. 60; Mismatches DΒ 1; ۲, Length

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Gaps

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RESULT 135 ABV68621 Human skin EST ABV68621; ABV68621 standard; cDNA; 11 21-OCT-2002 (first ВP

Human; skin; dermatological; vulnerary; antipsoriatic; antiseborrhaeic; immunosuppressive; antiinflammatory; cytostatic; SAGE; neurodermatitis; psoriasis; dermatitis; skin cancer; EST; expressed sequence tag; ss.

Homo sapiens.

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20-DEC-2001; 2001WO-EP015179

03-JAN-2001; 2001DE-01000127

(HENK ) HENKEL

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WPI; 2002-590638/63.

In vitro identification of a homeostasis and identifying .g. skin cancer. of skin-expressed genes, useful ying cosmetic or pharmaceutical a agents for determining

Disclosure; Page 203; 1345pp; German

in the skin of humans or animals by subjecting a mixture of genetically encoded factors from skin, to serial analysis of gene expression (SAGE) so as to identify skin-expressed genes and quantify their expression. (M1) is useful for identifying genes involved in skin homeostasis; to determine skin homeostasis and to test agent (A) that maintains or promotes skin homeostasis or that can be used for treating skin disorders, specifically namedate.

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RESULT 136
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                                                                                                                        The invention relates to in vitro identification (M1) of genes expressed in the skin of humans or animals by subjecting a mixture of genetically encoded factors from skin, to serial analysis of gene expression (SAGE) so as to identify skin-expressed genes and quantify their expression. (M1) is useful for identifying genes involved in skin homeostasis; to determine skin homeostasis and to test agent (A) that maintains or promotes skin homeostasis or that can be used for treating skin disorders, specifically neurodermatitis; sunburn; psoriasis; scoleroderma; ichthyosis; atopic dermatitis; acne; seborrhea; lupus erythematosus; rosacea; melanoma; basal cell carcinoma; and carcinoma or sarcoma of the skin. The present sequence is that of a human expressed sequence tag
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Human; skin; dermatological; vulnerary; antipsoriatic; antiseborrhaeic; immunosuppressive; antiinflammatory; cytostatic; SAGE; neurodermatitis; psoriasis; dermatitis; skin cancer; EST; expressed sequence tag; ss.
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                                                                                   Sequence
                                                                                                                                                                                                                                                                                               Disclosure; Page 69; 1345pp; German.
                                                                                                                                                                                                                                                                                                                            e.g.
                                                                                                                                                                                                                                                                                                                                      In vitro identification of skin-expressed genes, useful for determining homeostasis and identifying cosmetic or pharmaceutical agents against
                                                                                                                                                                                                                                                                                                                                                                                                            Petersohn D,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    03-JAN-2001; 2001DE-01000127.
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9; Conserv
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CTTCAGCAGC 1506
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Pred. No.
                                          Score 8.4; D
Pred. No. 60;
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RESULT 137
ARVESTIP
ARVESTIP
ID ABV652
XX ABV652
XX ABV652
XX TIMMUND
DT 21-OCT
XX HUMAN
XX HUMAN
XX HUMAN
XX HOMO 8
XX WO2002
PN WO2002
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XX WPI; 2
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ABV66909/c
ID ABV669
XX
AC ABV669
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AC ABV669
XX
DT 21-OCT
XX
DE Human
XX
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Best Local :
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 The invention relates to in vitro identification (M1) of genes expressed in the skin of humans or animals by subjecting a mixture of genetically encoded factors from skin, to serial analysis of gene expression (SAGE) so as to identify skin-expressed genes and quantify their expression.
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immunosuppressive; antiinflammatory; cytostatic; SAGE; neuro
psoriasis; dermatitis; skin cancer; EST; expressed sequence
                 Human skin EST 4695
                                                 21-OCT-2002
                                                                                     ABV66909;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                              (M1) is useful for identifying genes involved in skin homeostasis; determine skin homeostasis and to test agent (A) that maintains or promotes skin homeostasis or that can be used for treating skin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Disclosure; Page 108; 1345pp;
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RESULT 139
ABV71267/c
ID ABV712
XX ABV712
XX 21-OCT
XX 21-OCT
XX Human;
KW immuno
KW immuno
KW psoria
XX Homo s
XX Homo s
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Best Local :
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                                                                                                                                                                              Human; skin; dermatological; vulnerary; antipsoriatic; antiseborrhaeic; immunosuppressive; antiinflammatory; cytostatic; SAGE; neurodermatitis;
 03-JAN-2001; 2001DE-01000127
                                20-DEC-2001; 2001WO-EP015179
                                                                                                  WO200253774-A2
                                                                                                                                 Homo sapiens.
                                                                                                                                                                immunosuppressive; psoriasis; dermati
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human; skin; dermatological; vulnerary; antipsoriatic; antiseborrhaeic; immunosuppressive; antiinflammatory; cytostatic; SAGE; neurodermatitis; psoriasis; dermatitis; skin cancer; EST; expressed sequence tag; ss.
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Pred. No. 60;
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                                                                                                                                                                  EST;
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pharmaceutical agents against
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RESULT 140
ABV67181
ID ABV671
XX ABV671
XX ABV671
XX Human
XX Human
XX Human
XX Human
XX Homo 8
XX PO2002
XX Homo 8
XX WO2002
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                                                                                   e.g. skin cancer.
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                                                                                                                                                                                                                                                     Petersohn
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                                                                                                      disorders, specifically neurodermatitis; sunburn; psoriasis; scleroderma; ichthyosis; atopic dermatitis; ache; seborrhea; lupus erythematosus; rosacea; melanoma; basal cell carcinoma; and carcinoma or sarcoma of the
                                                                                                                                                                                                            The invention relates to in vitro identification (M1) of genes expressed in the skin of humans or animals by subjecting a mixture of genetically encoded factors from skin, to serial analysis of gene expression (SAGE) so as to identify skin-expressed genes and quantify their expression. (M1) is useful for identifying genes involved in skin homeostasis; to determine skin homeostasis and to test agent (A) that maintains or promotes skin homeostasis or that can be used for treating skin
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Cleavage_site
                                                                                                                                                                                                                                                                                                                                                                        of plants, animals, bacteria of subclones of the viral genome
                                                                                                                                                                                                                                                                                                                                                                                                  a recombinant viral genome, useful for of plants, animals, bacteria or viruses for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ₽P
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Pred. No. 6
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           chromosome; vaccine
                                                                                                                                                                                                                                                                                                                                                                             to assemble
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               C/DE-1 junction
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  0,
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The invention describes a method of directionally assembling a recombinant viral genome comprising ligating the subclones of the viral genome to assemble a recombinant viral genome, particularly coronavirus. For directionally assembling a recombinant viral genome. In particular, the method is useful for manipulating the genomes of higher plants and animals, as well as bacteria and viruses. In particular, the method is useful for the precise genetic manipulation of individual chromosomes in whole plants and animals and the construction of artificial chromosomes for gene therapy. The genomes produced are useful in preparing vaccines and expression vectors (e.g., TGE vectors and vaccines), which are useful in protocols involving vaccination, gene transfer and gene therapy. This sequence represents the interconnecting junction site C/DE-1 used in the

Example 7; Page 22; 42pp; English.

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RESULT 143
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XX ABT164
XX ABT164
XX Cytost
KW Cytost
KW Human
XX Cytost
KW Inflam
XX EP1262
XX Homo s
XX EP1262
XX O4-DEC
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                                                  The invention relates to a nucleic acid segment from the human neurokinin CC l receptor (TACR1) gene of 10-100 nucleotides comprising a fragment CC having a polymorphic site or a complement of the fragment. The TACR1 CC segment is useful for analysing a nucleic acid, by obtaining the nucleic acid from an individual, and determining the base occupying any one of the polymorphic sites in the segment. The nucleic acid is obtained from CC several individuals, and the base occupying one of the polymorphic sites is determined in each of the individuals, and further involves testing CC each of the individuals for the presence of a disease phenotype, and CC correlating the presence with the base. The TACR1 segment is useful for CC diagnosing and treating TACR1 ligand mediated diseases, such as asthma. CC The TACR1 segment is also useful in forensics, paternity testing, CC correlating polymorphisms with phenotypic traits, and genetic mapping of phenotypic traits. The TACR1 segment is useful in diagnosing and CC monitoring of diseases such as cancer, inflammation, heart disease, CC diseases of central nervous system, and susceptibility to infection to CC medicament for the treatment of the diseases. This polynucleotide sequence represents a polymorphic region of the human neurokinin 1 CC receptor (TACR1) gene of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Matches
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Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Novel nucleic acid segment from human neurokinin 1 receptor, including polymorphic sites for diagnosing and treating asthma, and in forensics paternity testing, and genetic mapping of the traits.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               genome
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Claim 1; Page 25; 27pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Cytostatic; antiasthmatic; antiinflammatory; cardiant; polymorphic human neurokinin 1 receptor; TACR1; disease phenotype; forensics; TACR1 ligand mediated disease; asthma; paternity testing; cancer; inflammation; heart disease; central nervous system; infection; ds
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ABT16435;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               assembly of the full length transmissible gastroenteritis virus (TGE) genome described in the method of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Affourtit JP,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               25-MAY-2001; 2001US-0293425P
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       20-MAR-2003 (first entry)
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90.0%;
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Sequence 11 BP; 3 A; 1 C; 3 G; 4 T; 0 U; 0 Other;

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Search completed: April 15, 2004, 16:35:36 Job time : 1 secs

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US-09-541-950B-19
US-08-541-950B-19
US-08-541-950B-22
US-09-083-756A-18
US-09-083-756A-19
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US-09-281-418-65
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AGCCAGACTTCAGCAGC	larity Conservat	, Application US/08 5821046 FORMATION: FORMATION: VINZENTION: VINZENTION: VIRAL P SEQUENCES: SEE: Banner & Witc SEE: Banner & Witc Companies ITYPE: Diskette, 3 ER: IBM PC compati ITYPE: Diskette, 3 ER: WordPerfect APPLICATION DATA: APPLICATION NUMBER: US/0 DATE: 03/19/5 PLICATION NUMBER: 07/9 MILIAMS, Ph.D., & WILLIAMS, PA.D., & WILLI		
AGCAC AGCAC	48. 76. ive	Jion US/08541  Jion US/08541  Jion US/08541  Jion US/08541  Jion US/0858  Jion Witcoff		10 10 10 10 10 10 10 10 11
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506 7	Sco. Pre	8541950B 8541950B IJ, Heaph (HIV) G Coff, Lt enter, 4 3.50 inc ible S/MS-DOS .1 08/541,9 960,370 1; Kathleen ,380 3777/5 70N: 00 17:	,	US-09- US-09- US-09- US-09- US-09- US-08- US-08- US-08- US-08- US-09- US-09-
	Mis Mis	1950B  Heaphy S, Dingw HF, Ltd. f, Ltd. er, 45th Floor e, 45th Floor 85-DOS 541,950B 541,950B 541,950B 777/57347 (MRC-	ALIGNMENTS	
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RESULT 2 US-08-541-950B-20 ; Sequence 20, Application US/08541950B

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 17, Application US/09083756A Patent No. 6114109
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: Karn J, C
TITLE OF INVENTION:
NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    FILING DATE: 03/19/93
ATTORNEY/AGENT INFORMATION:
NAME: Williams, Ph.D., Ke
REGISTRATION NUMBER: 34,
                                                                                                                                                                                                                                                                                                                                                                CORRESPONDENCE ADDRESS:
ADDRESSEE: Banner & Witcoff, Ltd.
STREET: One Financial Center, 45th Floor
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NUMBER OF SEQUENCES:
                                                                   CURRENT APPLICATION DATA
                                                                                                                                                                                                       COMPUTER READABLE FORM: MEDIUM TYPE: Diskett
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TOPOLOGY: linear MOLECULE TYPE: synthetic RNA
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TELECOMMUNICATION INFORMATION:
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STATE: Pu.
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STATE: FL.
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OTHER INFORMATION: N is 5-bromo-2'-deoxyuridine
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                                                                                                   COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS SOFTWARE: WordPerfect 6.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NAME/KEY: misc_feature
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TYPE: nucleic acid
STRANDEDNESS: single
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APPLICATION NUMBER: US/09/083,756A FILING DATE:
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One Financial Center, 45th Floor
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VENTION: VIRAL (HIV) GROWTH INHIBITION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Conservative
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                                                                                                                                                                                                       Diskette, 3.50 inch, 1.44 Mb storage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Gait MJ, Heaphy S, Dingwall C VIRAL (HIV) GROWTH INHIBITION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 48.0%;
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Pred. No. 2
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GENERAL INFORMATION:
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Best Local Similarity
                                                                                                                 TELEPHONE: (617) 345-9100
TELEFAX: (617) 345-9111
INFORMATION FOR SEQ ID NO: 20
SEQUENCE CHARACTERISTICS:
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                                                                                                                                                                                                                  ATTORNEY/AGENT INFORMATION:
NAME: Williams, Ph.D., Kathl
REGISTRATION NUMBER: 34,380
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SEQUENCE CHARACTERISTICS:
LENGTH: 17 bases
                                                                                                                                                                                     REFERENCE/DOCKET NUMBER: 3777/57347 (MRC-Ollax) TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                       FILING DATE:
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                            SOFTWARE: WordPerfect CURRENT APPLICATION DATA
                                                                                                                                                                                                                                                                                                                                                                                                                             COMPUTER READABLE FORM: MEDIUM TYPE: Diskette,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CORRESPONDENCE ADDRESS:
ADDRESSEE: Banner & Witcoff, Ltd.
STREET: One Financial Center, 45th Floor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: Karn J, Gait MJ, Heaphy S, Dingwall C TITLE OF INVENTION: VIRAL (HIV) GROWTH INHIBITION NUMBER OF SEQUENCES: 26
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TELECOMMUNICATION INFORMATION:
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APPLICATION NUMBER:
                                 MOLECULE TYPE:
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STATE: PL.
TO: 02111
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                                                                                                                                                                                                                                                                                                                                                                                          COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
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LOCATION: 8
OTHER INFORMATION: N is 2'-deoxythymidine
                                                                                                                                                                                                                                                                                                                                          APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NAME: Williams, Ph.D., Kathleen REGISTRATION NUMBER: 34,380
                                                                   STRANDEDNESS:
                                                                               TYPE: nucleic acid
                                                                                                                                                                                                                                                                                            APPLICATION NUMBER:
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   NAME/KEY:
                                                      TOPOLOGY:
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misc_feature
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Pred. No. 2.
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US-09-083-756A-20

48.0%; 76.5%;

Score 14.4; Pred. No. 2

DB 1;

Length 17;

OTHER INFORMATION:

N is 5-bromo-2'-deoxyuridine

VIRAL

(HIV) GROWTH INHIBITION

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US-09-083-756A-23; Sequence 23; Application US/09083756A; Patent No. 6114109; GENERAL INFORMATION: APPLICANT: Karn J, Gait MJ, Heaph;
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US-08-541-950B-23
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                                                                                             RESULT 6
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                                                                                                                                                                                                                                                         Query Match
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APPLICANT: Karn J
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TELEFAX: (617) 345-9111 INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NAME: Williams, Ph.D., Kathl
REGISTRATION NUMBER: 34,380
REFERENCE/DOCKET NUMBER: 37.
TELECOMMUNICATION INFORMATION:
TELEPHONE: (617) 345-9100
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        PRIOR APPLICATION DATA:
APPLICATION NUMBER: 07/90
FILING DATE: 03/19/93
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                SEQUENCE CHARACTERISTICS:
LENGTH: 18 bases
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SOFTWARE: WordPerfect
CURRENT APPLICATION DATA:
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CORRESPONDENCE ADDRESS:
ADDRESSEE: Banner & Witcoff, Ltd.
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                                                                                                                                                                                                                                                                                                                                                                              MOLECULE TYPE: synthetic RNA
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STATE: PL.
02111
PEAD/
                                                                                                                                                                                                                      Local Similarity
nes 13; Conserv
                                                                                                                                                                                                                                                                                                          OTHER INFORMATION: N is 4-thio-2'-deoxythymidine
                                                                                                                                                                                                                                                                                                                           NAME/KEY: misc_feature LOCATION: 8
                                                                                                                                                                                                                                                                                                                                                                                                             TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICATION NUMBER: UP
FILING DATE: 10/10/95
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS
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les 13; Conservative
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                                                                                                                                                                                   1490 AGCCAGACTTCAGCAGC 1506
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       Karn J, Gait MJ, Heaphy S, Dingwall C
                                                                                                                                                                                                                      Conservative
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                                                                                                                                                                                                                  48.0%; Score 14.4; 176.5%; Pred. No. 3.1 cive 2; Mismatches
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                                                                                                                US-09-325-601-1
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                                                                                                                            TITLE OF INVENTION: Methods and Kits for Discovery of RNA-Binding Compounds
FILE REFERENCE: 3950/81235
CURRENT APPLICATION NUMBER: US/09/325,601
CURRENT FILING DATE: 1999-06-03
NUMBER OF SEQ ID NOS: 53
SOFTWARE: Patentin Ver. 2.1
SEQ ID NO 1
LENGTH: 17
TYPE: RNA
ORGANISM: Human immunodeficiency virus
                                          Query Match
Best Local Similarity
Matches 13; Conserv
                                                                                                                                                                                                                                                                                                                                         Sequence 1, Application US/09325601 Patent No. 6573045
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INFORMATION FOR SEQ ID NO:
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APPLICANT: Prescott
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/541,950
FILING DATE:
ATTORNEY/AGENT INFORMATION:
NAME: Williams, Ph.D., Kathleen M.
REGISTRATION NUMBER: 34,380
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SOFTWARE: WordPerfect 6.1
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09,
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ADDRESSEE: Banner & Witcoff, Ltd.
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TELECOMMUNICATION INFORMATION:
TELEPHONE: (617) 345-9100
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MEDIUM TYPE: Diskett
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STATE: Pu.
STATE: 02111
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OTHER INFORMATION: N is 4-thio-2'-deoxythymidine
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NAME/KEY: misc_feature LOCATION: 8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS
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1490 AGCCAGACTTCAGCAGC 1506
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                                                  Conservative
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76.5%;
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                                                                   Score 13.8;
Pred. No. 3.
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RESULT 9
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APPLICANT: Karn J
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FILING DATE: 10/10/95
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 07/960,370
FILING DATE: 03/19/93
ATTORNEY/AGENT INFORMATION:
NAME: Williams, Ph.D., Kathleen
REGISTRATION NUMBER: 34,380
REGISTRATION NUMBER: 34,380
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INFORMATION FOR SEQ ID NO: 13:
SEQUENCE CHARACTERISTICS:
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette, 3.50 inch
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: WordPerfect 6.1
CURRENT APPLICATION DATA:
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MEDIUM TYPE: Diskett
COMPUTER: IBM PC com
OPERATING SYSTEM: PC
                                                                                                                                                       CORRESPONDENCE ADDRESS:
ADDRESSEE: Banner & Witcoff, Ltd.
STREET: One Financial Center, 45th Floor
                                                                                                                                                                                                                                 APPLICANT: Karn J, Gait MJ, Heaphy S, Dingwall C TITLE OF INVENTION: VIRAL (HIV) GROWTH INHIBITION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    MOLECULE TYPE: synthetic RNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SOFTWARE: WordPerfect 6.1
CURRENT APPLICATION DATA:
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CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                               NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TELECOMMUNICATION INFORMATION: TELEPHONE: (617) 345-9100
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TITLE OF INVENTION:
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STATE: Pu.
STATE: O2111
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STREET: CL.
STRY: Boston
WA
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STRANDEDNESS: single
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Similarity 76.5%;
                                                                                                                                            Boston
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ON: VIRAL (HIV) GROWTH INHIBITION
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                                                                        3.50 inch,
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Pred. No. 3
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                                                                       1.44 Mb storage
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CURRENT APPLICATION NUMBER: US/09/325,601; CURRENT FILING DATE: 1999-06-03; NUMBER OF SEQ ID NOS: 53; SOFTWARE: PatentIn Ver. 2.1; SEQ ID NO 3; SEQ ID NO 3; LENGTH: 18; TYPE: RNA; ORGANISM: Human immunodeficiency virus US-09-325-601-3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           US-09-325-601-3
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                                                                                                                                                                                                                                               RESULT 11
                                                                                                                   Patent No. 5821046
GENERAL INFORMATION:
GENERAL INFORMATION:
APPLICANT: Karn J, Gait MJ, Heaphy S, Dingwall C
TITLE OF INVENTION: VIRAL (HIV) GROWTH INHIBITION
NUMBER OF SEQUENCES: 26
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-- 'ocal Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TITLE OF INVENTION: Methods and Kits for Discovery of RNA-Binding Compounds FILE REFERENCE: 3950/81235
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: Karn
APPLICANT: Prescott
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LENGTH: 18 bases
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TELECOMMUNICATION INFORMATION:
TELEPHONE: (617) 345-9100
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ATTORNEY/AGENT INFORMATION:
NAME: Williams, Ph.D., Kathl
NAME: Williams, Ph.D., Kathl
NAME: 34,380
REGISTRATION NUMBER: 34,380
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PRIOR APPLICATION DATA:
APPLICATION NUMBER:
                                                                                  CORRESPONDENCE ADDRESS:
ADDRESSEE: Banner & Witcoff, Ltd.
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STREET: C...
STREET: MA
COMPUTER READABLE FORM:
   Cli.
STATE: MA
C2111
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STRANDEDNESS: single
TOPOLOGY: linear
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13; Conserv
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                                                                 One Financial Center, 45th Floor
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Pred. No. 3
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      DB
                                                                                                                                                                                                                                                                                                                                                                                                        DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                      Length 18;
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CURRENT APPLICATION DATA

WordPerfect 6.1 IBM PC compatible

Diskette, 3.50 inch, PC-DOS/MS-DOS

1.44 Mb storage

OPERATING SYSTEM: MEDIUM TYPE:

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
                              APPLICATION NUMBER: US/08/541,95
ETILING DATE: 10/10/95
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 07/960,370
PILING DATE: 03/19/93
ATTORNEY/AGENT INFORMATION:
NAME: W1111ams, Ph.D., Kathleen
REGISTRATION NUMBER: 34,380
REFERENCE/DOCKET NUMBER: 3777/57
TELECOMMUNICATION INFORMATION:
TELEPHONE: (617) 345-9100
                                                                                                                                                                                                                                                               STATE: MA
ZIP: 02111
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette, 3.50 inch,
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: WordPerfect 6.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          GENERAL INFORMATION:
TELEFAX: (617) 345-9111 INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TELEFAX: (617) 345-9111
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            tent No. 5821046
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ATTORNEY AGENT INFORMATION:
NAME: Williams, Ph.D., Kathl
REGISTRATION NUMBER: 34,380
                                                                                                                                                                                                                                                                                                                                                                                                                                         ADDRESSEE: Banner & Witcoff, Ltd.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      FILING DATE: 10/10/95
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 07
FILING DATE: 03/19/93
                                                                                                                                                                                                                                                CURRENT APPLICATION DATA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TITLE OF INVENTION:
NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    REFERENCE/DOCKET NUMBER: 3777/57347 (MRC-O11AX)
TELECOMMUNICATION INFORMATION:
TELEPHONE: (617) 345-9100
                                                                                                                                                                                                                                                                                                                                                                                       STREET: C...
STREET: C...
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Local 5.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TOPOLOGY: linear
MOLECULE TYPE: synthetic RNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             LOCATION: 9
OTHER INFORMATION: N is 2'-deoxythymidine
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NAME/KEY: misc_feature LOCATION: 9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ENGTH:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Application US/08541950B
                                                                                                                                                                                                                                                                                                                                                                                                                       One Financial Center, 45th Floor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           17 bases
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      42.7%; Score 12.8; Ilarity 76.5%; Pred. No. 5.6
Conservative 1; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Karn J, Gait MJ, Heaphy S, Dingwall C
TENTION: VIRAL (HIV) GROWTH INHIBITION
                                                                                                                                                                                                                                                                                                                         Diskette, 3.50 inch, 1.44 Mb storage
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                                                                                                              Kathleen M.
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                                                                          3777/57347 (MRC-Ollax)
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Best Local Similarity
                                                                                                                                                                                                                                                                             TELEFAX: (617) 345-9111 INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                         SOFTWARE: WordPerfect 6.1
CURRENT APPLICATION DATA;
APPLICATION NUMBER: US/08,
FILING DATE: 10/10/95
PRIOR APPLICATION DATA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette, 3.50 inch, 1.44 Mb storage
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                         ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CORRESPONDENCE ADDRESS:
ADDRESSEE: Banner & Witcoff, Ltd
                                                                                                                                                                    FEATURE:
                                                                                                                                                                                MOLECULE TYPE:
                                                                                                                                                                                                                                                               SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                             REFERENCE/DOCKET NUMBER: 37
TELECOMMUNICATION INFORMATION:
TELEPHONE: (617) 345-9100
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: Karn J, Gait MJ, Heaphy S, Dingwall CTITLE OF INVENTION: VIRAL (HIV) GROWTH INHIBITION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    MOLECULE TYPE:
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STATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NAME/KEY: misc_feature
LOCATION: 10
OTHER INFORMATION: N is 2'-deoxythymidine
                                                                                                                                                                                             TYPE: nucleic acid
STRANDEDNESS: single
TOPOLOGY: linear
                                                  Local
                                                                                                               LOCATION: 9 -
OTHER INFORMATION: N is 5-bromo-2'-deoxyuridine
                                                                                                                                          NAME/KEY: misc_feature
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            STREET:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                             NAME: Williams, Ph.D., REGISTRATION NUMBER: 3
                                                                                                                                                                                                                                                                                                                                                                                                              APPLICATION NUMBER: 0 FILING DATE: 03/19/93
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      LENGTH: 17 bases
                                                                                                                                                                                                                                             ENGTH:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1490 AGCCAGACTTCAGCAGC 1506
   1490 AGCCAGACTTCAGCAGC 1506
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         21, Application US/08541950B
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            02111
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                                                    Similarity
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                                                                                                                                                                                 synthetic RNA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     42.7%;
                                                  42.7%;
76.5%;
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                                                                                                                                                                                                                                                                                                                                                           34,380
                                                                                                                                                                                                                                                                                21:
                                                                                                                                                                                                                                                                                                                                                                              Kathleen M.
                                                                                                                                                                                                                                                                                                                                               3777/57347 (MRC-011AX)
                                   Score 12.8; D
Pred. No. 5.6;
1; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1; Mismatches
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Pred. No. 5.
                                                                  DB 1; Length 17;
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                                     Indels
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                                                                                                                                                                                                                                          Sequence 18, Application US/09083756A Patent No. 6114109
                                                                                                                                                                                                                                                                                                                                                                                                                          Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
Best Local Similarity
                                                                                                                                                                                                                             GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TELEFAX: (617) 345-9111
INFORMATION FOR SEQ ID NO: 22:
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette, 3.50 inch,
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                           APPLICANT: Karn J, Gait MJ, Heaphy S, Dingwall C TITLE OF INVENTION: VIRAL (HIV) GROWTH INHIBITION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    REFERENCE/DOCKET NUMBER: 37
TELECOMMUNICATION INFORMATION:
TELEPHONE: (617) 345-9100
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICATION NUMBER: US/08/541,9
FILING DATE: 10/10/95
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 07/960,370
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: Karn J, TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             MOLECULE TYPE: synthetic RNA FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SEQUENCE CHARACTERISTICS:
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MEDIUM TYPE: Diskett
                                                                                   ADDAL STREET: CALL BOSTON
                                                                                                                                                        CORRESPONDENCE ADDRESS:
                                                                                                                                                                            NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    FILING DATE: 03/19/93
ATTORNEY/AGENT INFORMATION:
NAME: Williams, Ph.D., Kathl
NAME: WILLIAMS, Ph.D., Kathl
REGISTRATION NUMBER: 34,380
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CURRENT APPLICATION DATA
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CORRESPONDENCE ADDRESS:
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STATE: DE 111
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STATE: PL.
TD. 02111
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TYPE: nucleic acid
STRANDEDNESS: singl
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          LOCATION: 10 OTHER INFORMATION: N is 5-bromo-2'-deoxyuridine
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  OPERATING SYSTEM:
SOFTWARE: WordPer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NAME/KEY: misc_feature
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                                                                                                                                                                                                                                                                                                                                                                                      1490 AGCCAGACTTCAGCAGC 1506
                                                                                                                                                                                                                                                                                                                                                                                                                          13;
                                                                                                                                                                                                                                                                                                                                                     1 AGCCAGAUUNGAGCAGC 17
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                                                                                                                        One Financial Center,
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                                                                                                                                                                                                                                                                                                                                                                                                                          Conservative
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                                                                                                                    Banner & Witcoff, Ltd.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    3: Diskette, 3.50 inch, 1.44 Mb storage
IBM PC compatible
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Banner & Witcoff, Ltd
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Gait MJ, Heaphy S, Dingwall C VIRAL (HIV) GROWTH INHIBITION
                                                                                                                                                                                                                                                                                                                                                                                                                                       42.7%;
76.5%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    PC-DOS/MS-DOS
                                                                                                                                                                          26
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          3777/57347 (MRC-011AX)
                                                                                                                                                                                                                                                                                                                                                                                                                                       Score 12.8;
Pred. No. 5
                                                                                                                                                                                                                                                                                                                                                                                                                          Mismatches
                                  1.44 Mb storage
                                                                                                                                                                                                                                                                                                                                                                                                                                                         Length 17;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 19,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Patent No. 6114109
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    GENERAL INFORMATION:
APPLICANT: Karn J
                                                                     TELEFAX: (617) 345-9111 INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TELEFAX: (617) 345-9111 INFORMATION FOR SEQ ID NO:
                                   SEQUENCE CHARACTERISTICS:
LENGTH: 17 bases
                                                                                                                    REFERENCE/DOCKET NUMBER: 37
TELECOMMUNICATION INFORMATION:
                                                                                                                                                    FILING DATE:
ATTORNEY/AGENT INFORMATION:
NAME: Williams, Ph.D., Kathleen M.
REGISTRATION NUMBER: 34,380
                                                                                                                                                                                                                                                                                            OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: WordPerfect 6.1
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NUMBER OF SEQUENCES: 2
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SEQUENCE CHARACTERISTICS:
LENGTH: 17 bases
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TELECOMMUNICATION INFORMATION: TELEPHONE: (617) 345-9100
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     PRIOR APPLICATION DATA:
APPLICATION NUMBER:
FILING DATE:
                                                                                                                                                                                                                                           PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                          COMPUTER READABLE FORM:
MEDIUM TYPE: Diskett
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               MOLECULE TYPE: synthetic RNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SOFTWARE: WordPerfect CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                            ADDxL.
STREET: U...
STREET: MA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NAME/KEY: misc_feature
LOCATION: 9
OTHER INFORMATION: N is 2'-deoxythymidine
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NAME: Williams, Ph.D., Kathl
REGISTRATION NUMBER: 34,380
REFERENCE/DOCKET NUMBER: 37
                                                                                                                                                                                                                                                             APPLICATION NUMBER: FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                              STATE: MA
ZIP: 02111
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       FILING
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICATION NUMBER:
TYPE: nucleic acid STRANDEDNESS: singl
                                                                                                   TELEPHONE:
                                                                                                                                                                                                                                                                                                                                              COMPUTER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                           APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TELEPHONE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ADDRESSEE: Banner & Witcoff, Ltd.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             POPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1490 AGCCAGACTTCAGCAGC 1506
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1 AGCCAGAUNUGAGCAGC 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       DATE
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                                                                                                                                                                                                                                                                                                                                                                                                                                                 One Financial Center, 45th Floor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Karn J,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WordPerfect 6.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 linear
                                                                                                                                                                                                                                                                                                                                            IBM PC compatible
                                                                                                      (617)
                                                                                                                                                                                                                                                                                                                                                              Diskette, 3.50 inch, 1.44 Mb
                                                                                                      345-9100
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Gait MJ, Heaphy S, Dingwall C VIRAL (HIV) GROWTH INHIBITION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         42.7%;
                                                                                                                                                                                                                           08/541,950
                                                                                                                                                                                                                                                                            US/09/083,756A
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Score 12.8; D
Pred. No. 5.6;
1; Mismatches
                                                                                                                                        3777/57347
                                                                                                                                          (MRC-011AX)
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                                                                                                                                                                                                                                                                                                                                                                storage
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TOPOLOGY: linear MOLECULE TYPE: synthetic RNA FEATURE:

NAME/KEY: misc\_feature

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RESULT 18
US-09-083-756A-22
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                                                                                                                                                                                                                                                                                                                                                                                                                                  ; NAME/KEY: misc_feature
LOCATION: 9
; OTHER INFORMATION: N is 5-bromo-2'-deoxyuridine
US-09-083-756A-21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       US-09-083-756A-21
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICATE:
PILING DATE:
ATTORNEY/AGENT INFORMATION:
NAME: Williams, Ph.D., Kathlee:
NAME: Will
                                                                                                                                                                                                                                                                                   Query Match 42. Best Local Similarity 76. Matches 13; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 21, Applicat Patent No. 6114109
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TELEFAX: (617) 345-9111 INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICATION NUMBER: US/09/083,756A
FILING DATE:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/541,950
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SOFTWARE: WordPerfect 6.1
CURRENT APPLICATION DATA:
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TITLE OF INVENTION: VIRAL (HIV) GROWTH INHIBITION
NUMBER OF SEQUENCES: 26
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette, 3.50 inch, 1.44 Mb
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       OLECULE TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Cli.
STATE: N.
STATE: 02111
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STRANDEDNESS: single
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                OPERATING SYSTEM:
SOFTWARE: WordPer
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OTHER INFORMATION: N is 2'-deoxythymidine
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Local Similarity
les 13; Conserv
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                                                                                                                                                                                                           1490 AGCCAGACTTCAGCAGC 1506
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                                                                                                                                               1 AGCCAGAUNUGAGCAGC 17
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One Financial Center, 45th Floor
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     IBM PC compatible
SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   synthetic RNA
                                                                                                                                                                                                                                                                                                                        76.5%;
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                                                                                                                                                                                                                                                                                                                        Score 12.8; D
Pred. No. 5.6;
                                                                                                                                                                                                                                                                                          Mismatches
                                                                                                                                                                                                                                                                                                                                                          DB 1; Length 17;
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RESULT 19
US-08-363-240A-47/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TELEPHONE: (617) 345-9100
TELEFAX: (617) 345-9111
INFORMATION FOR SEQ ID NO: 22
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APPLICANT: Karn J
                                                                                                                                                                                                                                                        GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           FILING DATE:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/54.
FILING DATE:
ATTORNEY/AGENT INFORMATION:
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: Karn J, Gait MJ, Heaphy S, Dingwall C TITLE OF INVENTION: VIRAL (HIV) GROWTH INHIBITION NUMBER OF SEQUENCES: 26
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SOFTWARE: WordPerfect 6.1
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COMPUTER READABLE FORM:

MEDIUM TYPE: Diskette, 3.50 inch,

COMPUTER: IBM PC compatible

OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                        APPLICANT: Pape, Michael TITLE OF INVENTION: METH
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SEQUENCE CHARACTERISTICS:
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                                                                                           CORRESPONDENCE ADDRESS:
                                                                                                                         TITLE OF INVENTION:
                                                                                                                                                                                                         APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      MOLECULE TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                REFERENCE/DOCKET NUMBER: 37
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                        APPLICANT:
                                                                                                                                                                                                                                         APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   C11.
STATE: ru.
C2111
                                        ADDRESSEE: Lyon & Lyon
STREET: 633 West Fifth
STREET: Suite 4700
                                                                                                                                                                                                                                                                                                                                                                                                                                                     Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       OTHER INFORMATION: N is 5-bromo-2'-deoxyuridine
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NAME/KEY: misc_feature
LOCATION: 10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CITY: Boston
COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     LOCATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NAME: Williams, Ph.D., REGISTRATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ADDRESSEE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COPOLOGY:
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                                                                                                                                                                                                                                                                                                                                                                       1 AGCCAGAUUNGAGCAGC 17
                                                                                                                                                                                                                                                                          5705388
           Los Angeles
California
                                                                                                                                                                                                                                                                                      Application US/08363240A
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One Financial Center, 45th Floor
                                                                                                                                                                                                     McSwiggen, James
Bisgaier, Charles
                                                                                                                                                                                                                                                                                                                                                                                                                                        Conservative
                                                                                                                                                                                                                                         Couture, Larry
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    synthetic RNA
                                                                                                                                                           PREVENTION,
                                                                                                                                                                        METHOD AND REAGENT FOR
                                                                                                                         PROGRESSION OF VASCULAR
                                                                                                                                                                                                                                                                                                                                                                                                                                                       42.7%;
76.5%;
                                                                                                            1243
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                                                            Street
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                                                                                                                                                                                                                                                                                                                                                                                                                                                       Score 12.8;
Pred. No. 5.
                                                                                                                       N AND REGRESSION R DISEASES
                                                                                                                                                                                                                                                                                                                                                                                                                                        Mismatches
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Patent No. 5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Matches 13;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
Best Local Similarity 86.7%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                GENERAL INFORMATION:
CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: Petry, Douglas A.
REGISTRATION NUMBER: 35,321
REFERENCE/DOCKET NUMBER: 8765
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICATION NUMBER: US
FILING DATE: December
PRIOR APPLICATION DATA:
APPLICATION NUMBER:
FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ZIP: 90071

COMPUTER READABLE FORM:
MEDIUM TYPE: Storage
MEDIUM TYPE: Storage
                                                                                       CURRENT APPLICATION DATA:
APPLICATION NUMBER: US
FILING DATE:
                                                                                                                                                                          COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SEQUENCE CHARACTERISTICS:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ATTORNEY/AGENT INFORMATION:
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                                                                                                                                                                                                                                                                                              CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                               NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                            TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TELEFAX: 67-3510
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              LENGTH: 15 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                            COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TELEPHONE: (213) 955-0440
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COMPUTER: IBM Compatible OPERATING SYSTEM: IBM P.C. SOFTWARE: Word Perfect 5.1
                                                                                                                                                                                                                                                   STREET: 340 K
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NAME: Warburg, Richard REGISTRATION NUMBER: 32. REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                           COUNTRY: UZIP: 07110
                                                                                                                                                                                                                                                                                 ADDRESSEE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1488 GAAGCCAGACTTCAG 1502
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5567809
                                                                                                                                                                                                                       New Jersey
                                                                                                                                                                                                                                                                                    340 Kingsland
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Begovich, Aun ...
Begovich, Aun ...
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                                                                                                                                                                                                                                                                                 Hoffmann-La Roche Inc
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        December 23, 1994
                                                                                                                                                                                                                                                                                                                                                                                                                               Raymond
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                                                                                                       US/08/050,073
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Pred. No. 7
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                                                                                                                                                                                                                                                                                                                                          US-09-508-753B-25/c
                                                                                                                                                                                                                                                                                                                                                             RESULT 22
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APPLICANT: Indue, Takakazu

APPLICANT: Indue, Takakazu

TITLE OF INVENTION: Method of Amplifying DNA Fragment, Apparatus for Amplifying DNA Fr
TITLE OF INVENTION: agment, Method of Assaying Microorganisms, Method of Analyzing Mic
TITLE OF INVENTION: nisms and Method of Assaying Contaminant

FILE REFERENCE: 9982-7

CURRENT APPLICATION NUMBER: US/09/281,418

CURRENT FILING DATE: 1999-03-30

EARLIER APPLICATION NUMBER: JP/1998/87651

EARLIER APPLICATION NUMBER: JP/1998/87651

EARLIER APPLICATION NUMBER: JP/1999/69694

EARLIER APPLICATION NUMBER: JP/1999/69694

EARLIER FILING DATE: 1999-03-16
                                                                                                                                                                        APPLICANT: AKIRA SHIMAMOTO
APPLICANT: Yasuhiro FURUICI
APPLICANT: Yuko SHIBATA
APPLICANT: Hiroko FUNAKI
APPLICANT: Eiji OHARA
APPLICANT: Eiji OHARA
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NUMBER OF SEQ ID NOS:
SEQ ID NO 25
                                                                                                                                                                                                                                                                              GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                 Sequence 25, Application US/09508753B Patent No. 6544736
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SEQ ID NO 65
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
                                 CURRENT APPLICATION NUMBER: US/09/508,753B
CURRENT FILING DATE: 2000-06-16
PRIOR APPLICATION NUMBER: JP 9/270324
PRIOR FILING DATE: 1997-09-18
                                                                                                                    APPLICANT: Masanori WATAHIKI
TITLE OF INVENTION: Method for Synthesizing
FILE REFERENCE: 00162/HG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NUMBER OF SEQ ID NOS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TELEFAX: (510) 814-2977 INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TYPE: DNA
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      OTHER INFORMATION: Primer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     LENGTH: 12
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10; Conserv
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Yuko SHIBAJA
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Pred. No. 16;
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Pred. No. 8.7;
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RESULT 24
US-09-721-777-18
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Best Local Similarity
"---hes 9; Conserv:
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US-09-263-790-36
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         CURRENT APPLICATION NUMBER: US/09/721,777
CURRENT FILING DATE: 2000-11-27
NUMBER OF SEQ ID NOS: 20
SOFTWARE: FASTSEQ for Windows Version 4.0
SEQ ID NO 18
LENGTH: 10
TYPE: DNA
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: Nirmal Kumar PATRA et al.

APPLICANT: Nirmal Kumar PATRA et al.

TITLE OF INVENTION: JAL PALLAVI, WATER LOGGING TOLERANT CYMBOPOGON WINTERIANUS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SOFTWARE:
SEQ ID NO 36
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Best Local Similarity
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CURRENT FILING DATE: 1999-03-05
NUMBER OF SEQ ID NOS: 38
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TYPE: DNA
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                                                                                                                                                                                                                                                                                                                                            APPLICANT: Khanuja, Suman Preet Singh
APPLICANT: Kumar, Sushil
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ORGANISM: Artificial FEATURE:
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FEATURE:
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INVENTION: Mint Plant Named Saksham
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                                                                                                                                                                                                                                                                                                        Shasany, Ajic ,
Dhawan, Sunita
                                                                                                                                                                                                                                                                                                                                                                                                                Application US/09721777
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    PatentIn version 3.0
                                                                                                                                                                                                    Singh, Anil Kumar
Patra, Nirmal Kumar
Bahl, Janak Raj
                                                                                                                                                                                                                                                       Naqvi, Ali Arif
Dhawan, Om Parkash
                                                                                                                                                                                                                                                                                           Darokar, Mahendra Pandurang
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100.0%; Pred. No. 16;
tive 0; Mismatches
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o. 16;
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Query Match
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                                                                                                          US-08-545-253A-20
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Best Local Similarity 90.0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     GENERAL INFORMATION:
                                                                                                                                                                                                     TELEX: 575102
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                           REGISTRATION NUMBER: 31,665
REFERENCE/DOCKET NUMBER: 50
TELECOMMUNICATION INFORMATION:
TELEPHONE: (919) 881-3140
                                                                                                                                                                                       SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                              CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: Sibley, Kenneth D.
                                                                                                                                                                                                                                                                                                                                                                                            COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CORRESPONDENCE ADDRESS:
ADDRESSEE: Kenneth D. Sibley
STREET: Post Office Drawer 34009
                                                                                                                    TOPOLOGY: 15 MOLECULE TYPE:
                                                                                                                                                                                                                                                                                                                                                                    SOFTWARE: PatentIn Release #1.0, Version CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TITLE OF INVENTION:
TITLE OF INVENTION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                  STATE: N
COUNTRY:
                                                                                                                                              STRANDEDNESS: single
                                                                                                                                                                                                                       TELEPHONE: (919) 881-3175
                                                                                                                                                                                                                                                                                                                                                       APPLICATION NUMBER:
                                                                                                                                                                          ENGTH:
                         1488 GAAGCCAGAC 1497
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   E. George Kuhlman
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  U.S.A.
                                                   Conservative
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                                                                                                                                     linear
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90.0%;
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                                                               Score 8.4;
Pred. No.
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Pred. No.
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RESULT 26 US-08-719-337-20 ; Sequence 20, Application US/08719337 ; Patent No. 6054634

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RESULT 27
US-09-255-432-6
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CLASSIFICATION: 047

CLASSIFICATION DATA:
APPLICATION NUMBER: 08/184,567

FILING DATE: 21-JAN-1994

ATTORNEY/AGENT INFORMATION:
NAME: Sibley, Kenneth D.
REGISTRATION NUMBER: 31,665

REGISTRATION NUMBER: 5051-2

TELECOMMUNICATION:
"THEORMATION:
"THEORMATI
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Best Local Similarity
Matches 9; Conserva
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 6, Application US/09255432
Patent No. 6258537
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ENERAL INFORMATION
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
                                                                                     COMPUTER READABLE FORM:
MEDIUM TYPE: Diskett
                                                                                                                                                                                                                                                                                                                                                                             CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                              NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                               TITLE OF INVENTION:
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CITY: No. 6054634th Carolina
                                                                                                                                                                                                                   CITY: Greenville
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COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TELEPHONE: (919) 881-3175
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICATION NUMBER: US/0 FILING DATE: 25-SEP-1996
COMPUTER: IBM compatible OPERATING SYSTEM: MS Dos; Windows SOFTWARE: WordPerfect 6.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              STRANDEDNESS: single
                                                                                                                                                              ZIP:
                                                                                                                                                                                                                                                                       ADDRESSEE: Judy C. Jarecki-Black, Ph.D.

ADDRESSEE: Dority & Manning, P.A.

STREET: 700 E. No. 6258537th Street, Suite 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TOPOLOGY:
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Post Office Drawer 34009
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                                                                                         Diskette,
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                                                                                         3.50 inch, 1.44 Mb storage
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Pred. No. ;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Patent No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
                                                                                PRIOR APPLICATION DATA:
APPLICATION NUMBER: A:
FILING DATE: 21 June
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                        GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CURRENT APPLICATION DATA:
APPLICATION NUMBER: US
FILING DATE: Filed Her
                                                                SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                COMPUTER: IBM Compatible w/
OPERATING SYSTEM: Microsoft
SOFTWARE: Microsoft word 97
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                    COMPUTER READABLE FORM: MEDIUM TYPE: Diskett
                                                                                                                                                                                                                                                                                                                                                 CORRESPONDENCE ADDRESS:
ADDRESSEE: D. Peter Hochberg Co., L.P.A.
STREET: The Baker Building - Sixth Floor 1940 East 6th Street
                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: William Mitchell Molyneux TITLE OF INVENTION: Mosquito and/or Flea Control NUMBER OF SEQUENCES: 20
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ANTI-SENSE: NO
ORIGINAL SOURCE: Operon Technologies (Alemeda,
IMMEDIATE SOURCE: Operon Technologies
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DESCRIPTION:
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ATTORNEY/AGENT INFORMATION:
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                                                                                                                                                                                                                                                                                                                                      STREET: The STATE: Ohio
                                                                                                                                                                APPLICATION NUMBER: US
FILING DATE: June 19,
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REFERENCE/DOCKET NUMBER:
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               STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                       COUNTRY:
                                                                                                                                                  CLASSIFICATION:
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TYPE: Nucleic Acid
                                               ENGTH:
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o. 6337071
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                             Nucleic Acid
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                                                                                                                                                                                                                                                                                                                       U.S.A.
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Linear
                                                                                                                                                                                                                                                                      Diskette,
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                                                                                                                                                    300
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)N: No. 6258537 Applicable
                                                                                                                   AU PO 0605
                                                                                                                                                                  US/08/878,835A
9, 1997
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 P44,
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                                                                                                 1996
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                                                                                                                                                                                                                                                                      3.5 inch,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CXU-291
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                                                                                                                                                                                                                                      Windows
                                                                                                                                                                                                                                                      Pentium Processor
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                                                                                                                                                                                                                                                                      1.44 MByte storage
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FILE REFERENCE: 00162/HG
CURRENT APPLICATION NUMBER: US/09/508,753B
CURRENT FILING DATE: 2000-06-16
PRIOR APPLICATION NUMBER: JP 9/270324
PRIOR FILING DATE: 1997-09-18
NUMBER OF SEQ ID NOS: 472
SEQ ID NO 63
LENGTH: 10
TYPE: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                               RESULT 30
US-09-508-753B-63/c
                                                 ; OTHER INFORMATION: Description of Artificial Sequence: Primer US-09-508-753B-63
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CURRENT APPLICATION NUMBER: US/09/508,7538
CURRENT FILING DATE: 200-06-16
PRIOR APPLICATION NUMBER: JP 9/270324
PRIOR FILING DATE: 1997-09-18
NUMBER OF SEQ ID NOS: 472
SEQ ID NO 28
LENGTH: 10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
Best Local Similarity
Matches 9; Conserv
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Best Local Similarity
Matches 9; Conserv
Query Match
Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                         Description US/09508753B Patent No. 6544736
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GENERAL INFORMATION:
                                                                                                                                                                                                                                                            APPLICANT: Masanori WATAHIKI
TITLE OF INVENTION: Method for Synthesizing
                                                                                                                                                                                                                                                                                                                APPLICANT:
                                                                                                                                                                                                                                                                                                                                               APPLICANT: Akira SHIMAMOTO
APPLICANT: Yasuhiro FURUICHI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: Akira SHIMAMOTO APPLICANT: Yasuhiro FURUI
                                                                                                                                                                                                                                                                                             APPLICANT:
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                                                                                ORGANISM: Artificial Sequence FEATURE:
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                                                                                                                                                                                                                                                                                                                Hiroko FUNAKI
                                                                                                                                                                                                                                                                                                                               Yuko SHIBATA
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Yuko SHIBATA
                                                                                                                                                                                                                                                                                                  Eiji OHARA
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Pred. No. 20;
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Pred. No. 20;
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                 DB 1;
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                                                                                                                                                                                                                                                              cDNA from mRNA sample
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                                                                                                                        RESULT 32
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                                                                       Sequence 6, Application US/09758073 Patent No. 6610487
                                                                                                                                                                                                                                             Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 110, Application US/08894454 Patent No. 6544784
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                                                         GENERAL INFORMATION:
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           APPLICANT: Keinath, TITLE OF INVENTION: TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                           SEQUENCE CHARACTERISTICS:
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MEDIUM TYPE: Diskette
COMPUTER: IBM Compatil
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ADDRESSEE: The Webb Law Firm
STREET: 700 Koppers Building, 436 Seventh Avenue
                                                                                                                                                                                                                                                                                                                                                                                                                                          TELECOMMUNICATION INFORMATION:
TELEPHONE: 412-471-8815
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICATION NUMBER: PCT/EP
FILING DATE: 19-FEB-1996
APPLICATION NUMBER: 952003
FILING DATE: 17-FEB-1995
APPLICATION NUMBER: 952019
FILING DATE: 14-UUL-1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: VAN DEN VEN, W.J.M.
APPLICANT: SCHOENMAKERS, H.F.P.M.
TITLE OF INVENTION: MULTIPLE-TUMOR ABERRENT GROWTH
TITLE OF INVENTION: GENES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ATTORNEY/AGENT INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                               REFERENCE/DOCKET NUMBER: 703
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NAME: Johnson, Barbara E
REGISTRATION NUMBER: 31,
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CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICATION NUMBER:
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SOFTWARE: FastSEQ for Windows Version
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Plants Using a Polymerase Chain Reaction Assay
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Pred. No. 20;
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TUMBER OF SEQUENCES:

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US-08-173-489C-342/c
; Sequence 342, Application US/08173489C
; Patent No. 5861244
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                                                                                                                                                                                            Patent No. 5861244
GENERAL INFORMATION:
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Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                             Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TELEFAX: (864) 233-7342 INFORMATION FOR SEQ ID NO: (
                                                                                                                            APPLICANT: WANG, C. APPLICANT: HEPBURN, A TITLE OF INVENTION: CITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                        OTHER INFORMATION: Commercially Available Primer PUBLICATION INFORMATION: No. 6610487 Applicable
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TELECOMMUNICATION INFORMATION: TELEPHONE: (864) 271-1592
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                FILING DATE: 16-MAR-1998
ATTORNEY/AGENT INFORMATION:
NAME: Judy C. Jarecki-Black, Ph.D.
REGISTRATION NUMBER: P44,170
                                                                                          NUMBER OF SEQUENCES: 3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TOPOLOGY: Li
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APPLICATION NUMBER:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ADDRESSEE: Judy C. Jarecki-Black, Ph.D.
ADDRESSEE: Dority & Manning, P.A.
STREET: 700 E. No. 6610487th Street, Suite
CITY: Greenville
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OPERATING SYSTEM:
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                              STATE: NEW YORK
                                                              STREET:
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                COUNTRY:
                                                                             ADDRESSEE:
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Oligonucleotide Primer
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                                                                                                                              GENETIC SEQUENCE ASSAY USING DNA TRIPLE-STRAND FORMATION.
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Pred. No. 20;
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Query Match
Best Local Similarity
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                                                                                                      ; OTHER INFORMATION: Symthetic oligonucleotide primer. US-09-862-847-15
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                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 15, Applica Patent No. 6593111 GENERAL INFORMATION:
                                                                                                                                                                                                                   SEQ ID NO 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
                                                                                                                                                                                                                                                   CURRENT APPLICATION NUMBER: US/09/862,847
CURRENT FILING DATE: 2001-05-21
PRIOR APPLICATION NUMBER: US 60/206,537
PRIOR FILING DATE: 2007-05-21
PRIOR APPLICATION NUMBER: US 60/285,320
PRIOR FILING DATE: 2001-04-20
NUMBER OF SEQ ID NOS: 24
                                                                                                                                                                                                                                        SOFTWARE: PatentIn version 3.1
                                                                                                                                                                                                                                                                                                                                                                                  TITLE OF INVENTION: DIRECTION ASSEMBLY OF LARGE VIRAL GENOMES AND CHROMOSOMES FILE REFERENCE: 5470.270
                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: Baric, Ralph S. APPLICANT: Boyd, Youn:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TELEFAX: (attorney) (: INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                 TYPE: DNA
                                                                                                                                                               ORGANISM: Artificial Sequence
                                                                                                                                                                                                    ENGTH: 11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ANTI-SENSE: no PUBLICATION INFORMATION:
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MOLECULE TYPE: other nucleic acid

DESCRIPTION: third strand derived from P. cepacea

DESCRIPTION: 16s region in Seq ID No. 5861244341
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LENGTH: 11 bases
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICATION NUMBER: US 07/968,436 FILING DATE: 29 OCT 1992 ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5 inch, 1.44Mb storage
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                RELEVANT RESIDUES IN SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NAME: Handelman, Joseph H.
REGISTRATION NUMBER: 26,179
REFERENCE/DOCKET NUMBER: U9518-6
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1484 CCAAGAAGCC 1493
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                                                    Score 8.4; I
Pred. No. 21;
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Pred. No. 21;
                                     Mismatches
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RESULT 36
US-09-041-675-19/c
; Sequence 19, Application US/09041675A
; Patent No. 6100032
; Patent No. 6100032
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US-08-859-954-95/c
                                                                                                    GENERAL INFORMATION:
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APPLICANT: Kern, Scott
APPLICANT: Zawel, Leigh
APPLICANT: Dai, Jia Le
APPLICANT: Vogelstein, Bert
APPLICANT: Kinzler, Kenneth
TITLE OF INVENTION: Human SMAD3 and SMAD4 are
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MOLECULE TYPE: otne
DESCRIPTION: /des
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NAME: Paul, Thomas D.
REGISTRATION NUMBER: 32,714
REFERENCE/DOCKET NUMBER: D-59
TELECOMMUNICATION INFORMATION:
TELEPHONE: 713/651-5325
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TELEFAX: 713/651-5246
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    GENERAL INFORM
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: IBM PC COMPATIBLE
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/859,954
                                                                                                                                                                                                                                                                                                                                                                               ANTI-SENSE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CLASSIFICATION:
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: Hardin, Paul E.
TITLE OF INVENTION: Design and Optimized Primer Library for
TITLE OF INVENTION: Gene Sequencing and Method Thereof
NUMBER OF SEQUENCES: 566
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      STATE: Terri
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                                                                                                                                                                                                                                                                                          Local Similarity es 8; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                              STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TYPE: nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COUNTRY: U.S.A. ZIP: 77010-3095
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ENGTH: 8 base pairs
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1301 McKinney, Suite 5100
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GENERAL INFORMATION:
APPLICANT: Kern, Scott
APPLICANT: Dai, Jia Le
APPLICANT: Dai, Jia Le
APPLICANT: Dai, Jia Le
APPLICANT: Vogelatein, Bert
APPLICANT: Vogelatein, Bert
APPLICANT: Vogelatein, Bert
APPLICANT: Winzler, Kenneth
TITLE OF INVENTION: Human SMAD3 and SMAD4 are
TITLE OF INVENTION: sequence-specific transcription activators
FILE REFERENCE: 01107.74098
CURRENT APPLICATION NUMBER: US/09/041,675A
CURRENT APPLICATION NUMBER: US/09/041,675A
CURRENT FILING DATE: 1998-03-13
NUMBER OF SEQ ID NOS: 27
SOFTWARE: FASTSEQ for Windows Version 3.0
SEQ ID NO 24
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US-09-989-789-455/c
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Best Local Similarity
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GENERAL INFORMATION:
APPLICANT: LIU, Qiang
TITLE OF INVENTION: POSITION DEPENDENT RECOGNITION OF GNN NUCLEOTIDE
TITLE OF INVENTION: TRIPLETS BY ZINC FINGERS
FILE REFERENCE: 8325-0011.20 / S11-US2
CURRENT APPLICATION NUMBER: US/09/989,789
CURRENT FILING DATE: 2002-03-25
NUMBER OF SEQ ID NOS: 4085
SOFTWARE: Patentin Ver. 2.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CURRENT APPLICATION NUMBER: US/09/041,675A
CURRENT FILING DATE: 1998-03-13
NUMBER OF SEQ ID NOS: 27
SOFTWARE: FASTSEQ for Windows Version 3.0
SEQ ID NO 19
                                                                                                                                                                                      Sequence 455, Application US/09989789 Patent No. 6588746
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 LENGTH: 8
TYPE: DNA
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    OTHER INFORMATION:
OTHER INFORMATION:
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OTHER INFORMATION: synthetic random oligonucleotides selected
OTHER INFORMATION: binding to human SMAD3 or human SMAD4
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TYPE: DNA
ORGAN_SM: Artificial Sequence
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100.0%; Pr
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; Pred. No.
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APPLICANT: LIU, Qiang
TITLE OF INVENTION: POSITION DEPENDENT RECOGNITION OF GNN NUCLEOTIDE
TITLE OF INVENTION: TRIPLETS BY ZINC FINGERS
FILE REFERENCE: 8325-0011.20 / S11-US2
CURRENT APPLICATION NUMBER: US/09/989,789
CURRENT FILING DATE: 2002-03-25
NUMBER OF SEQ ID NOS: 4085
SOFTWARE: PATENTIN Ver. 2.0
SEQ ID NO 456
LENGTH: 9
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Best Local Similarity
Matches 8; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TYPE: DNA
                   ZIE: 90071-2066
COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette,
MEDIUM TYPE: storage
                                                                                                                                                                                                                             TITLE OF INVENTION: TITLE OF INVENTION:
                                                                                                                                                                                                                                           APPLICANT: Jerry W. Shay
APPLICANT: Woodring E. Wright
APPLICANT: Elizabeth Blackburn
TITLE OF INVENTION: THERAPY AND DIAGNOSIS OF CONDITIONS
TITLE OF INVENTION: RELATED TO TELOMERE LENGTH AND/OR
                                                                                                                                                                                           CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                             NUMBER OF SEQUENCES:
                                                                                                                                       STREET: 633 West Fifth Street
STREET: Suite 4700
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      / Match 26.7%; So Local Similarity 100.0%; les 8; Conservative 0;
                                                                                                            STATE:
                                                                                         COUNTRY: U.S.A.
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E: California
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nilarity 100.0%;
Conservative 0;
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Compatible
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57
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0; Mismatches
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                                       1.44 Mb
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.2e+02;
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Query Match
Best Local Similarity
Matches 8; Conserv:
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SOFTWARE: PatentIn Ver. 2.0
SEQ ID NO 4
LENGTH: 10
TYPE: DNA
ORGANISM: Mycobacterium tuberculosis
US-08-997-897-4
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Best Local Similarity
Matches 8; Conserv:
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US-09-156-836B-4
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                                                                                                                                                                                                                                                                                                                                                                              Sequence 4, Application US/08997897C
Patent No. 6114514
GENERAL INFORMATION:
APPLICANT: SRIVASTAVA, RANJANA
APPLICANT: SRIVASTAVA, BRAHM SHANKER
APPLICANT: SRIVASTAVA, BRAHM SHANKER
TITLE OF INVENTION: MYCOBACTERIUM TUBERCULOSIS SPECIFIC DNA FRAGMENT
FILE REFERENCE: U011469-7
CURRENT APPLICATION NUMBER: US/08/997,897C
CURRENT FILING DATE: 1997-12-24
                                            Sequence 4, Application US/09156836B Patent No. 6242585 GENERAL INFORMATION:
 APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TELEX: 67-3510
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                              APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICATION NUMBER: 07/882,438
FILING DATE: May 13, 1992
APPLICATION NUMBER: 08/018,766
FILING DATE: March 24, 1993
ATTORNEY/ACENT IMPORMATION:
NAME: DATE: DATE: DATE: THORNATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                REFERENCE/DOCKET NUMBER: 20
TELECOMMUNICATION INFORMATION:
TELEPHONE: (213) 489-1600
TELEPAX: (213) 955-0440
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CURRENT APPLICATION DATA: APPLICATION NUMBER: US
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NAME: Warburg, Richard J.
REGISTRATION NUMBER: 32,327
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            OPERATING SYSTEM:
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Srivastava, Ranjana
Kumar, Deepak
Srivastava, Brahm Sh
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Pred. No.
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5. 23;
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RESULT 43
US-08-464-011B-9/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Matches
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PRIOR FILING DATE: 1997-12-24
NUMBER OF SEQ ID NOS: 7
SOFTWARE: Description of the control of the contro
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TYPE: DNA
ORGANISM: Mycobacterium tuberculosis
                                                                                                             TELEX: 67-3510
INFORMATION FOR SEQ ID NO: 9:
                                                                                                                                                                                                                                                                                           APPLICATION NUMBER: 07/882,438
FILING DATE: May 13, 1992
APPLICATION NUMBER: 08/038,766
FILING DATE: March 24, 1993
APPLICATION NUMBER: 08/060,952
FILING DATE: May 13, 1993
ATTORNEY/AGENT INFORMATION:
NAME: Warburg, Richard J.
REGISTRATION NUMBER: 32,327
REFERENCE/DOCKET NUMBER: 202/045
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICATION NUMBER: US/08/464,011B
FILING DATE: 05-Jun-1995
CLASSIFICATION: <Unknown>
PRIOR APPLICATION DATA:
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TITLE OF INVENTION: THERAPY AND DIAGNOSIS OF CONDITIONS
RELATED TO TELOMERE LENGTH AND/OR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1486 AAGAAGCC 1493
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CORRESPONDENCE ADDRESS:
                                                                SEQUENCE CHARACTERISTICS:
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LENGTH: 10
TYPE: nucle
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CITY: Los Angeles
STATE: California
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STREET: 633 West Fifth Street
                                                                                                                                                                                                                      TELEPHONE:
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                                                                                                                                                                                  TELEFAX: (213) 955-0440
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                                                                                                                                                                                                                          (213) 489-1600
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Pred. No. 23;
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Search completed: April 15, 2004, 16:36:47 Job time: 0.001 secs

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                                                                                                                                      ; OTHER INFORMATION: Description of Artificial Sequence:Synthetic oligonucleotide US-09-336-946B-15
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US-09-336-946B-15
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                                                                                                                                                                                                                 CURRENT APPLICATION NUMBER: US/09/336,946B; CURRENT FILING DATE: 1999-06-21; PRIOR APPLICATION NUMBER: 60/095229; PRIOR FILING DATE: 1998-08-04; NUMBER OF SEQ ID NOS: 74; SOFTWARE: Microsoft Office 97; SEQ ID NO 15; LENGTH: 10
                                                              Query Match
Best Local Similarity
Matches 8; Conserv
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Best Local (
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APPLICANT: Bryan, Gregory
APPLICANT: E. I. du Pont de Nemours and Company
TITLE OF INVENTION: A Pi-ta GENE CONFERRING DISEASE RESISTANCE TO PLANTS
FILE REFERENCE: BB-1136
                                                                                                                                                                         TYPE: DNA ORGANISM: Artificial Sequence
                                                                                                                                                                        FEATURE:
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nes 8; Conserv
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2 AGCCAGAC 9
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                                                                Conservative
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100.0%; Pred. No.
                                                            26.7%; Score 8; DB 1
100.0%; Pred. No. 23;
tive 0; Mismatches
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Sequence 65, Appl
Sequence 67, Appl
Sequence 1799, Ap
Sequence 1800, Ap
Sequence 40, Appl
Sequence 41, Appl
Sequence 1451, Ap
Sequence 1451, Ap
Sequence 11451, Ap
Sequence 11451, Ap
Sequence 1145, Ap
Sequence 1137, Ap
Sequence 11, Appl
Sequence 11, Appl
Sequence 11, Appl
Sequence 14, Appl
Sequence 14, Appl
Sequence 14, Appl
Sequence 1647, Appl
Sequence 1647, Appl
Sequence 1647, Appl
Sequence 1651, Appl
Sequence 1651, Appl
Sequence 1107, Appl
Sequence 1107, Appl
Sequence 1107, Appl
Sequence 292, App
Sequence 1107, Appl
Sequence 1107, Appl
Sequence 1551, Appl
Sequence 1551, Appl
Sequence 455, Appl
                                                                                                                      APPLICANT: Brett P. Monia
APPLICANT: Susan M. Freier
APPLICANT: Scott Cooper
APPLICANT: Scott Cooper
TITLE OF INVENTION: ANTISENSE MODULATION OF FIBROBLAST GROWTH FACTOR RECEPTOR 2 EXPRES
FILE REFERENCE: RTS-0250
CURRENT APPLICATION NUMBER: US/09/954,556
CURRENT FILING DATE: 2001-09-14
NUMBER OF SEQ ID NOS: 108
SEQ ID NO 66
LENGTH: 20
TYPE: DNA
ORGANISM: Artificial Sequence
FEATURE:
OTHER INFORMATION: Antisense Oligonucleotide
US-09-954-556-66
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US-09-954-556-65/c
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Matches 20
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Matches 20
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Publication No. US20030078219A1
GENERAL INFORMATION:
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SEQ ID NO 65
LENGTH: 20
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TITLE OF INVENTION: ANTISENSE MODULATION OF FIBROBLAST GROWTH FACTOR RECEPTOR 2 EXPRES FILE REFERENCE: RTS-0250 CURRENT APPLICATION NUMBER: US/09/954,556 CURRENT FILING DATE: 2001-09-14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: Brett P. Monia
APPLICANT: Susan M. Freier
APPLICANT: Scott Cooper
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 1484 CCAAGAAGCCAGACTTCAGC 1503
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                                                       66.7%;
ilarity 100.0%;
Conservative (
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100.0%; Pr
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US-09-989-994-56
US-10-113-877-5
US-10-339-161-6
US-10-277-494-147
US-08-463-404-9
US-10-033-145-185
US-10-033-145-1815
US-10-133-030-3
US-10-358-818-3
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Pred. No.
                                                                        Score 20; DB 1;
Pred. No. 0.62;
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US-10-307-005-1800
US-11-307-005-1800
US-11-004-551-43
US-09-504-231A-1451
US-09-504-231A-1435
US-09-274-553D-1435
US-09-274-553D-1435
US-09-274-553D-1435
US-09-274-553D-145-1327
US-09-823-887C-17
US-10-106-799-13
US-10-106-799-13
US-10-106-799-13
US-10-106-799-13
US-10-106-799-13
US-10-106-799-14
US-09-822-250-14
US-09-822-250-14
US-09-823-377-14
US-10-331-15-1647
US-10-331-15-1647
US-10-331-15-1611
US-10-331-1627-292
US-10-330-627-1077
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US-10-342-615-110
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Post-processing: Minimum Match 0%
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Listing first 44 summaries

Database

rnpb.seq:

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

Minimum

Total number of hits satisfying chosen parameters:

86

43 seqs, 507 residues

Gapop 10.0 , IDENTITY\_NUC

Gapext 0.5

Minimum DB Maximum DB

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length: 8 length: 50

Sequence: Title:

Perfect score:

us-09-954-556-3 30

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GenCore version (c) 1993 - 2004

5.1.6 Compugen Ltd.

April 15, 2004, 16:38:43; Search time 0.001 Seconds (without alignments) 30.420 Million cell updates/sec

Scoring table:

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APPLICANT: Howard B. Gamper
APPLICANT: Howard B. Gamper
APPLICANT: Michael C. Rice
APPLICANT: Michael C. Rice
APPLICANT: Jungsup Kim
TITLE OF INVENTION: Targeted Chromosomal Genomic Alterations in Plants
TITLE OF INVENTION: Using Modified Single Stranded Oligonucleotides
FILE REFERENCE: Napro/009 PCT
CURRENT APPLICATION NUMBER: US/10/307,005
CURRENT FILING DATE: 2002-11-26
PRIOR APPLICATION NUMBER: US/10/307/05
PRIOR APPLICATION NUMBER: US 60/208,538
PRIOR APPLICATION NUMBER: US 60/208,538
PRIOR APPLICATION NUMBER: US 60/244,989
PRIOR FILING DATE: 2000-06-01
PRIOR FILING DATE: 2000-10-30
PRIOR FILING DATE: 2000-10-30
PRIOR FILING DATE: 2000-10-30
PRIOR FILING DATE: 2001-05-01
PRIOR FILING DATE: 2001-05-01
PRIOR FILING DATE: 2001-05-01
PRIOR FILING DATE: 2001-03-27
PRIOR FILING DATE: 2001-03-27
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; Sequence 67, Application US/09954556

; Publication No. US20030078219A1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CURRENT APPLICATION NUMBER: US/09/954,556
CURRENT FILING DATE: 2001-09-14
NUMBER OF SEQ ID NOS: 108
SEQ ID NO 67
LENGTH: 20
TYPE: DNA
ORGANISM: Artificial Sequence
                                                                                      Matches
                                                                                                                                 Query Match
                                                                                                                                                                                                                                                                      SEQ ID NO 1799
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 1799, Application US/10307005
Publication No. US20030236208A1
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                                                                                                                                                                                                                                                                                         NUMBER OF SEQ ID NOS: 2717
SOFTWARE: Friedman macro N
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: Susan M. Freier
APPLICANT: Scott Cooper
TITLE OF INVENTION: ANTISENSE MODULATION OF FIBROBLAST GROWTH FACTOR RECEPTOR 2
FILE REFERENCE: RTS-0250
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: Brett P. Monia
APPLICANT: Susan M. Frei
APPLICANT: Scott Cooper
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: University of Delaware
                                                                                                                                                                                                  ORGANISM: Solanum tuberosum
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                                                                                                                                                                                                                                               ENGTH: 17
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ses 20; Conservative
                                       1485 CAAGAAGCCAGACTTCA 1501
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CAAGAAGCTAAACTTCA 17
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                                                                                                         46.0%;
88.2%;
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100.0%; Pred. No. 0.
Live 0; Mismatches
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Pred. No. 4
                                                                                    Mismatches
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0.62;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 40, Application US/10004551
Publication No. US20030004310A1
GENERAL INFORMATION:
APPLICANT: SHIMEETS, RICHARD A
APPLICANT: FERNANDES, ELMA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SOFTWARE: Friedman macro Napro4
SEQ ID NO 1800
LENGTH: 17
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                                                                                                                                                                                                                                                                                       SEQ ID NO 40
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APPLICANT: Howard B. Gamper
APPLICANT: Michael C. Rice
APPLICANT: Jungs:p Kim
ITITLE OF INVENTION: Targeted Chromosomal Genomic Alterations in Plants
ITITLE OF INVENTION: Using Modified Single Stranded Oligonucleotides
FILE REFERENCE: Napro/009 PCT
CURRENT APPLICATION NUMBER: US/10/307,005
CURRENT FILING DATE: 2002-11-26
PRIOR APPLICATION NUMBER: PCT/US01/17672
PRIOR FILING DATE: 2001-06-01
                                                                                                                                                                                                                                                                                                        CURRENT APPLICATION NUMBER: US/10/004,551
CURRENT FILING DATE: 2001-12-05
PRIOR APPLICATION NUMBER: 09/635,949
PRIOR FILING DATE: 2000-08-10
NUMBER OF SEQ ID NOS: 110
SOFTWARE: Patentin Ver: 2.1
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PRIOR FILING DATE: 2001-03-27
NUMBER OF SEQ ID NOS: 2717
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PRIOR APPLICATION NUMBER: US 60/208,538
PRIOR FILING DATE: 2000-06-01
PRIOR APPLICATION NUMBER: US 60/244,989
PRIOR FILING DATE: 2000-10-30
                                                                                                                                                                                                                                                                                                                                                                                                                                         FILE REFERENCE: 15966-559
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                                                                                                                                                                                                                                                                                                                                                                                                                                                          TITLE OF INVENTION: POLYNUCLEOTIDES AND POLYPEPTIDES ENCODED THEREBY
                                                                                                                                                                                                                           ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                  TYPE: DNA
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les 15; Conserv
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                                      1481 CGACCAAGAAGCCAGAC 1497
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88.2%;
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Pred. No. 4.8;
0; Mismatches
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Pred. No. 5
                                                                                 Mismatches
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US-09-504-231A-1451
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                                                                                                                                                                                                                                                                                                      PRIOR APPLICATION NUMBER: 09/274,553
PRIOR FILING DATE: 1999-03-23
PRIOR PELICATION NUMBER: 09/257,608
PRIOR FILING DATE: 1999-02-24
PRIOR APPLICATION NUMBER: 60/100,842
PRIOR APPLICATION NUMBER: 60/100,842
PRIOR FILING DATE: 1998-09-18
PRIOR APPLICATION NUMBER: 60/083,217
PRIOR FILING DATE: 1998-04-27
NUMBER OF SEQ ID NOS: 3242
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US-09-504-231A-1451/c
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PRIOR FILING DATE: 2000-08-10
NUMBER OF SEQ ID NOS: 110
SOFTWARE: Patentin Ver. 2.1
SEQ ID NO 43
                                                                                                                                                                                                                                                 SOFTWARE: PatentIn version 3.0
SEQ ID NO 1451
LENGTH: 14
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                                                                                             Query Match
Best Local Similarity
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Matches 15; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                       FILE REFERENCE: rpi 247/282
CURRENT APPLICATION NUMBER: US/09/504,231A
CURRENT APPLICATION NUMBER: US/09/504,231A
CURRENT FILING DATE: 2000-02-15
PRIOR APPLICATION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TITLE OF INVENTION: POLYNUCLEOTIDES AND POLYPEPTIDES ENCODED THEREBY FILE REFERENCE: 15966-559
CURRENT APPLICATION NUMBER: US/10/004,551
CURRENT FILING DATE: 2001-12-05
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: SHIMKETS, APPLICANT: FERNANDE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TITLE OF INVENTION: ENZYMATIC NUCLEIC TITLE OF INVENTION: HEPATITIS C VIRUS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     OTHER INFORMATION: Description of Artificial Sequence: PCR PRIMER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ORGANISM: Artificial Sequence FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TYPE: DNA
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                                                                                                                                                                                                               ORGANISM: Artificial Sequence
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0. US20020013458A1
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                                      CTTCAGCAGCCA 1508
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               McSwiggen, James
Roberts, Beth
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  FERNANDES,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Macejak, Dennis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Pavco, Pamela
                                                                             Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              2001-12-05
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                                                                                               40.0%; Score 12; DB 1; 100:0%; Pred. No. 7.3;
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Pred. No. 5
                                                                             Mismatches
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                                                                                                                 Length 14;
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                                                                             Indels
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US-09-504-231A-1435/c
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; OTHER INFORMATION: Description of Artificial Sequence: US-09-504-231A-1435
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                                                                                                                          SOFTWARE: Patentin
SEQ ID NO 1435
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    GENERAL INFORMATION
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Best Local Similarity
Matches 12; Conserv
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APPLICANT: Pavco, Pamela
APPLICANT: Macejak, Dennis
TITLE OF INVENTION: ENZYMATIC NUCLEIC ACID TREATMENT OF DISEASES OR CONDITIONS RELATED
TITLE OF INVENTION: HEPATITIS C VIRUS INFECTION
FILE REFERENCE: Tpi 247/282
CURRENT APPLICATION NUMBER: US/09/274,553D
CURRENT FILING DATE: 1999-03-23
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PRIOR APPLICATION NUMBER: 60/100,842
PRIOR FILING DATE: 1998-09-18
                                                                                                                                                                                                                                                                                                                                                              FILE REFERENCE: rpi 247/282
CURRENT APPLICATION NUMBER: US/09/504,231A
CURRENT FILING DATE: 2000-02-15
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                                                                                                                                                                       NUMBER OF SEQ ID NOS: 3242
                                                                                                                                                                                           PRIOR APPLICATION NUMBER: 60/083,217 PRIOR FILING DATE: 1998-04-27
                                                                                                                                                                                                                                   PRIOR FILING DATE: 1999-02-24
PRIOR APPLICATION NUMBER: 60/100,842
PRIOR FILING DATE: 1998-09-18
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                                                                                                                                                                                                                                                                                                  PRIOR FILING DATE: 1999-03-23
PRIOR APPLICATION NUMBER: 09/257,608
                                                                                                                                                                                                                                                                                                                                          PRIOR APPLICATION NUMBER: 09/274,553
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APPLICANT: McSwig
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                                                            ORGANISM: Artificial Sequence
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ITLE OF INVENTION: ENZYMATIC NUCLEIC ACID TREATMENT OF DISEASES OR CONDITIONS RELATED
TILE OF INVENTION: HEPATITIS C VIRUS INFECTION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  FEATURE:
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Pred. No.
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                        Nucleic Acid Target
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Sequence 1327, Application US/10033145

Publication No. US20020151515A1

GENERAL INFORMATION:
APPLICANT: GENZYME CORPORATION
APPLICANT: ROBERTS, BRUCE
APPLICANT: SHANKARA, SRINIVAS
TITLE OF INVENTION: PREPARATION AND USE OF SUPERIOR VACCINES
FILE REFERENCE: GA0201C
CURRENT APPLICATION NUMBER: US/10/033,145
CURRENT FILING DATE: 2001-11-05
PRIOR APPLICATION NUMBER: PCT/US99/13800
PRIOR FILING DATE: 1999-06-18
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                                                              ; ORGANISM: Homo sapiens US-10-033-145-1327
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PRIOR FILING DATE: 1999-02-24
PRIOR APPLICATION NUMBER: 60/100,842
PRIOR FILING DATE: 1998-09-18
PRIOR APPLICATION NUMBER: 60/083,217
PRIOR FILING DATE: 1998-04-27
NUMBER OF SEQ ID NOS: 3148
                                                                                                                                         NUMBER OF SEQ ID NOS: 2137
SOFTWARE: PatentIn version 3.0
SEQ ID NO 1327
Query Match
Best Local Similarity
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Best Local
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SEQ ID NO 1435
LENGTH: 14
TYPE: RNA
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APPLICANT: McSwiggen, James
APPLICANT: Roberts, Beth
APPLICANT: Roberts, Beth
APPLICANT: Pavco, Pamela
APPLICANT: Macejak, Dennis
APPLICANT: Macejak, Dennis
TITLE OF INVENTION: ENZWATIC NUCLEIC ACID TREATMENT OF DISEASES OR CONDITIONS RELATE
TITLE OF INVENTION: ENZWATIC VIRUS INFECTION
FILE REFERENCE: Tpi 247/282
CURRENT APPLICATION NUMBER: US/09/274,553D
CURRENT APPLICATION NUMBER: US/09/274,553D
CURRENT FILING DATE: 1999-03-23
                                                                                                         TYPE: DNA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                OTHER INFORMATION: Description of Artificial Sequence: Nucleic
                                                                                                                            LENGTH: 10
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85.7%; Pred. No. 11;
tive 0; Mismatches
33.3%;
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Pred. No. 11;
  Score 10;
Pred. No.
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  DB 1;
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                 Length 10;
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Query Match
Best Local Similarity
Thes 10; Conserve
RESULT 15
US-10-240-580-10/c
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APPLICANT: Lal, Lakhvir
APPLICANT: Alal, Lakhvir
TITLE OF INVENTION: Cloning of No. US20030180723Ale
TITLE OF INVENTION: Dormancy in the Apical Buds of
FILE REFERENCE: HO53916.0001US0
CURRENT APPLICATION NUMBER: US/09/823,887C
CURRENT FILING DATE: 2002-04-23
NUMBER OF SEQ ID NOS: 33
SOFTWARE: PatentIn version 3.1
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US-10-106-799-13
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Publication No. US20030140379A1
GENERAL INFORMATION:
APPLICANT: COUNCIL of Scientific and Industrial Research
TITLE OF INVENTION: No. US20030140379A1el DNA sequence in plants Caragana jubata with
TITLE OF INVENTION: method thereof
FILE REFERENCE: US 673
                                                                                                                                                                                                                                                                                                                  CURRENT APPLICATION NUMBER: US/10/106,799
CURRENT FILING DATE: 2002-10-31
NUMBER OF SEQ ID NOS: 32
SOFTWARE: Patentin version 3.1
SEQ ID NO 13
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Matches 10; Conserv
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LENGTH: 13
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ORGANISM: Artificial Sequence
FEATURE:
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Pred. No.
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Pred. No.
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he Apical Buds of Tea (Camellia Sinensis L. (O.) Kur
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13;
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Sequence 10, Application US/10240580 Publication No. US20030180716A1

GENERAL INFORMATION:

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RESULT 16
US-08-935-377-14
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Publication No. US20030133917A1
GENERAL INFORMATION:
APPLICANT: Zauderer, Maurice
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match 31.3%;
Best Local Similarity 90.9%;
Matches 10; Conservative
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SEQ ID NO 10
LENGTH: 12
TYPE: DNA
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TITLE OF INVENTION: METHOD AND APPARATUS FOR MICROORGANISM DISCRIMINATION, METHOD OF
TITLE OF INVENTION: DATABASE FOR MICROORGANISM DISCRIMINATION, AND MICROORGANISM DISCRIMINATION OF RECORDED TITLE REPERENCE: 9982-24
CURRENT PILING DATE: 2002-09-30
PRIOR APPLICATION NUMBER: PCT/JP01/02516
PRIOR PILING DATE: 2001-03-27
PRIOR APPLICATION NUMBER: JP 2000-99482
PRIOR FILING DATE: 2000-03-31
NUMBER OF SEQ ID NOS: 46
                                                                                                                                                                    CLASSIFICATION: 424
ATTORNEY/AGENT INFORMATION:
NAME: Steffe, Eric K
REGISTRATION NUMBER: 36,688
REGERENCE/DOCKET NUMBER: 1821
TELECOMMUNICATION INFORMATION:
TELEPHONE: (202) 371-2600
                                                                                                      TELEFAX: (202) 371-254
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       OTHER INFORMATION: Primer
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                                                                                                                                                                                                                                                                                                                                                                          COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOCTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ZIP: 20005
COMPUTER READABLE FORM:
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CORRESPONDENCE ADDRESS:
MOLECULE TYPE:
                                  STRANDEDNESS:
                                                            TYPE: nucleic acid
                                                                                                                                                                                                                                                                                                                              APPLICATION NUMBER: US/0 FILING DATE: 22-SEP-1997
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CITY: Washington
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ADDRESSEE: Sterne, Kessler, Goldstein & Fox P.L.L.C
STREET: 1100 New York Avenue, N.W., Suite 600
                    TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                MEDIUM TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                1484 CCAAGAAGCCA 1494
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                                                                                                                                                    (202) 371-2540
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                                     single
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                                                                                                                                                                                                                     1821.0010000/EKS/CMB
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Pred. No. 15;
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Query Match

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Score 9;

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1;

Length 10;

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RESULT 19
US-10-044-674-86/c
US-10-040-674-86/c
; Sequence 86, Application US/10044674
; Publication No. US20030175710A1
; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ; LENGTH: 10
TYPE: DNA
; ORGANISM: Synthetic construct
US-09-822-250-14
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Best Local Similarity
Tatches 9; Conserv:
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; Sequence 1647, Application US/10033145
; Publication No. US20020151515A1
; GENERAL INFORMATION:
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Best Local Similarity
Wardhes 9; Conserva
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APPLICANT: Zauderer, Maurice
TITLE OF INVENTION: Methods for Producing Recombinant Libraries in Vaccinia Virus
FILE REFERENCE: 1821.0010001
FILE REFERENCE: 1821.0010001
CURRENT FILING DATE: 2001-04-02
PRIOR APPLICATION NUMBER: US 08/935,377
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               PRIOR FILING DATE: 1997-09-22
NUMBER OF SEQ ID NOS: 37
SOFTWARE: PatentIn version 3.0
SEQ ID NO 14
                                                                                                                                                                                                                                                                                              NUMBER OF SEQ ID NOS: 2137
SOFTWARE: PatentIn version 3.0
SEQ ID NO 1647
LENGTH: 10
                                                                                                                                                                                                                                                                                                                                                                               TITLE OF INVENTION: PREPARATION AND USE OF SUPERIOR VACCINES FILE REFERENCE: GAOZOIC CURRENT PILING DATE: 2001-11-05
PRIOR APPLICATION NUMBER: US/10/033,145
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 14, Application US/09822250 Patent No. US20020018785A1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: GENZYME CORPORATION
                                                                                                                                                                                                                                                                                  TYPE: DNA
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100.0%; Pred. No.
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Best Local Similarity
7-1-ches 9; Conserv
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SEQ ID NO 86
LENGTH: 10
                                                                                                                REGISTRATION NUMBER: P44,170
REFERENCE/DOCKET NUMBER: CXU-
TELECOMMUNICATION INFORMATION:
TELEPHONE: (864) 271-1592
TELEFAX: (864) 233-7342
INFORMATION FOR SEQ ID NO: 6:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 6, Application US/09758073
Patent No. US20010024785A1
GENERAL INFORMATION:
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TITLE OF INVENTION: HAPLTYPES OF THE TNFRSF11B GENE
FILE REFERENCE TWERSF11B MWH-0001US (CIP)
CURRENT APPLICATION NUMBER: US/10/044,674
CURRENT FILING DATE: 2002-01-09
PRIOR APPLICATION NUMBER: PCT/US00/18803
PRIOR PLICATION DATE: 2000-07-10
NUMBER OF SEQ ID NOS: 94
                                                                                                                                                                                                                                                                                                                                                          COMPUTER: IBM compatible
OPERATING SYSTEM: MS Dos; W
SOFTWARE: WordDerfect 6.1
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/7
FILING DATE: Filed Herewith
CLASSIFICATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TYPE: DNA
                                                                                                                                                                                                                                                      PRIOR APPLICATION DATA:
APPLICATION NUMBER: 60/078,103
FILING DATE: 16-MAR-1998
ATTORNEY/AGENT INFORMATION:
NAME: Judy C. Jarecki-Black, Ph.D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: Keinath, et al.
TITLE OF INVENTION: Method of Diagnosing Gummy Stem Blight in
TITLE OF INVENTION: Plants Using a Polymerase Chain Reaction Assay
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    COMPUTER READABLE FORM:
MEDIUM TYPE: Diskett
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NUMBER OF SEQUENCES: 1
                  TOPOLOGY: Linear
MOLECULE TYPE: Other Nucleic Acid
                                         LENGTH: 10 Pairs
TYPE: Nucleic Acid
STRANDEDNESS: Single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ADDRESSEE: Judy C. Jarecki-Black, Ph.D.
ADDRESSEE: Dority & Manning, P.A.
STREET: 700 E. No. US20010024785Alth Street,
CITY: Greenville
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       1496 ACTTCAGCA 1504
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Stephens, J. Claiborne
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Oligonucleotide Primer
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100.0%; Pr
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ER: CXU-291
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; Pred. No. 14;
0; Mismatches
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Query Match
Best Local Similarity
Tatches 9; Conserve
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Best Local Similarity
Tarches 9; Conserve
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; OTHER INFORMATION: Exon/intron of TORB
US-09-772-105-77
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SEQ ID NO 77
LENGTH: 10
                                                                                                                                                                                 Sequence 1651, Application US/10033145
Publication No. US20023151515A1
GENERAL INFORMATION:
APPLICANT: GENZYME CORPORATION
APPLICANT: ROBERTS, BRUCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                GENERAL INFORMATION:
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Patent No. US20010029015A1
SOFTWARE: PatentIn version 3.0 SEQ ID NO 1651
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APPLICANT: Breakefield, Xandra O.
TITLE OF INVENTION: TORSIN, TORSIN-RELATED GENES, AND
TITLE OF INVENTION: METHODS OF DETECTING NEURONAL DISEASES
                                                 CURRENT APPLICATION NUMBER: US/10/033,145
CURRENT FILING DATE: 2001-11-05
PRIOR APPLICATION NUMBER: PCT/US99/13800
PRIOR FILING DATE: 1999-06-18
                                                                                                                             APPLICANT: SHANKARA, SRINIVAS
TITLE OF INVENTION: PREPARATION AND USE OF SUPERIOR VACCINES
FILE REFERENCE: GA0201C
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        PRIOR APPLICATION NUMBER: US PRIOR FILING DATE: 1997-06-19 NUMBER OF SEQ ID NOS: 90
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PRIOR APPLICATION NUMBER: US 09/218,363
PRIOR FILING DATE: 1998-12-22
PRIOR APPLICATION NUMBER: US 09/099,454
PRIOR FILING DATE: 1998-06-18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CURRENT APPLICATION NUMBER: US/09/772,105
CURRENT FILING DATE: 2001-01-26
                                     NUMBER OF SEQ ID NOS: 2137
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ORGANISM: Unknown
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TYPE: DNA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         OTHER INFORMATION: COT PUBLICATION INFORMATION:
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Operon Technologies
MB: No. US20010024785A1 Applicable
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Pred. No. 17;
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PRIOR APPLICATION NUMBER: US 09/448,480
PRIOR FILING DATE: 1999-11-24
PRIOR DATE: PASTSEQ for Windows Version 4.0
SEQ ID NO 292
LENGTH: 10
TYPE: DNA
OPERANTE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           RESULT 24
US-10-330-627-292
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US-10-033-145-1651
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                                                                                                                                 ; ORGANISM: Homo sapiens
US-10-330-627-292
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APPLICANT: Kinzler, Kenneth W
APPLICANT: Kinzler, Kenneth W
APPLICANT: Vogelstein, Bert
TITLE OF INVENTION: Human Transcriptomes
FILE REFERENCE: 001107.00319
CURRENT APPLICATION NUMBER: US/10/330,627
CURRENT FILING DATE: 2002-12-30
PRIOR APPLICATION NUMBER: US 09/448,480
PRIOR FILING DATE: 1999-11-24
NUMBER OF SEQ ID NOS: 1564
                                                       Query Match 28.0%;
Best Local Similarity 90.0%;
Matches 9; Conservative
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Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 292, Applic Publication No. US20 GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                          APPLICANT: Velculescu, Victor E.
APPLICANT: Kinzler, Kenneth W
APPLICANT: Vogelstein, Bert
TITLE OF INVENTION: Human Transcriptomes
FILE REFERENCE: 001107.00319
                                                                                                                                                                                                                                                                                                    CURRENT APPLICATION NUMBER: US/10/330,627
CURRENT FILING DATE: 2002-12-30
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TYPE: DI
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les 9; Conservative
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1499 TCAGCAGCCA 1508
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b. US20030175771A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                      Application US/10330627 o. US20030175771A1
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90.0%;
                                                                          Score 8.4; DB 1; Length 10; Pred. No. 17;
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Pred. No. 17;
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Pred. No. 1
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Query Match
Best Local Similarity
Watches 9; Conserva
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US-10-330-627-1077/c
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US-10-352-615-110
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SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 1077
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 110, Application US/10352615 Publication No. US20030190285A1 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CURRENT APPLICATION NUMBER: US/10/330,627
CURRENT FILING DATE: 2002-12-30
PRIOR APPLICATION NUMBER: US 09/448,480
PRIOR FILING DATE: 1999-11-24
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: Vogelstein, Bert
TITLE OF INVENTION: Human Tra
FILE REFERENCE: 001107.00319
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: Velculescu, Victor E.
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                                                                                                     APPLICATION NUMBER: US/08/894,454
FILING DATE: 15-AUG-1997
APPLICATION NUMBER: PCT/EP/00716
FILING DATE: 19-FEB-1996
APPLICATION NUMBER: 95200390.3
FILING DATE: 17-FEB-1995
APPLICATION NUMBER: 95201951.1
FILING DATE: 14-JUL-1995
APPLICATION NUMBER: 95201951.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COUNTER READABLE FORM:
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
COMPUTER: IBM Compatible
COMPUTER: DOS
COMPUTER: DOS
COMPUTER: COMPATIBLE
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APPLICATION NUMBER: US/10/352,615
FILING DATE: 28-Jan-2003
CLASSIFICATION: 424
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ADDRESSEE: The Webb
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TITLE OF INVENTION: MULTIPLE-TUMOR ABERRENT GROWTH
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REFERENCE/DOCKET NUMBER: 702-971100
                                      NAME: Johnson, Barbara E
REGISTRATION NUMBER: 31,1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SOFTWARE: FastSEQ
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Building,
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Pred. No.
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RESULT 27
US-09-862-847-15/c
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LENGTH: 11
TYPE: DNA
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Matches
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Best Local :
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APPLICANT: Webb, Suzin M.
APPLICANT: Affourtit, Jason P.
APPLICANT: Affourtit, Jason P.
TITLE OF INVENTION: GENETIC POLYMORPHISMS IN THE HUMAN NEUROKININ 1 RECEPTOR GENE AND TITLE OF INVENTION: USES IN DIAGNOSIS AND TREATMENT OF DISEASES
FILE REFERENCE: PC10461AGPR
CURRENT APPLICATION NUMBER: US/10/146,354A
CURRENT FILING DATE: 2002-08-15
PRIOR APPLICATION NUMBER: 60/293,425
PRIOR FILING DATE: 2001-05-25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             PRIOR APPLICATION NUMBER: US 60/285,320 PRIOR FILING DATE: 2001-04-20 NUMBER OF SEQ ID NOS: 24
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CURRENT APPLICATION NUMBER: US/09/862,847
CURRENT FILING DATE: 2001-05-21
PRIOR APPLICATION NUMBER: US 60/206,537
PRIOR FILING DATE: 2000-05-21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SOFTWARE: PatentIn version 3.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TITLE OF INVENTION: DIRECTION ASSEMBLY OF FILE REFERENCE: 5470.270
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: Baric, Ralph S. APPLICANT: Boyd, Yount
                                                                                                                                                                                                          APPLICANT: Pfizer Inc
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TELEX: <Unknown>
INFORMATION FOR SEQ ID NO: 110:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              OTHER INFORMATION: Synthetic oligonucleotide primer
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TELEPHONE: 412-471-8815
TELEFAX: 412-471-4094
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                                                                                                                                                                                                                                                                                                                                                                                                                                                          Similarity
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TYPE: nucleic acid
STRANDEDNESS: single
TOPOLOGY: linear
                                                                                                                                                                   Seymour, Albert B.
Nelson, Darcy L.
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Pred. No. 17;
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Pred. No. 19;
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Query Match
Best Local Similarity
Matches 10; Conserv
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; ORGANISM: Homo sapiens
US-10-146-354A-16
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                                                              US-09-989-789-455
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                                                                                                                                                        CURRENT APPLICATION NUMBER: US/09/989,789
CURRENT FILING DATE: 2002-03-25
NUMBER OF SEQ ID NOS: 4085
SOFTWARE: PatentIn Ver. 2.0
SEQ ID NO 455
LENGTH: 9
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                                                                                                                                                                                                                                                                                                                   GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                Sequence 455, Application US/09989789 Patent No. US20020063379A1
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LENGTH: 20
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Best Local Similarity
Matches 8; Conserv
                               Query Match
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CURRENT FILING DATE: 2001-09-14
NUMBER OF SEQ ID NOS: 108
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                                                                                                                                                                                                                                                   APPLICANT: LIU, Qiang
TITLE OF INVENTION: POSITION DEPENDENT RECOGNITION OF
TITLE OF INVENTION: TRIPLETS BY ZINC FINGERS
FILE REFERENCE: 8325-C011.20 / S11-US2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TITLE OF INVENTION: ANTISENSE MODULATION OF FIBROBLAST GROWTH FACTOR RECEPTOR 2 FILE REFERENCE: RTS-0250
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APPLICANT: Susan M. Freier
APPLICANT: Scott Cooper
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                                                                            OTHER INFORMATION: Description of Artificial Sequence: example target OTHER INFORMATION: DNA
                                                                                                                       ORGANISM: Artificial Sequence
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                                                                                                                                            TYPE: DNA
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90.0%;
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76.9%;
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              Score 8; 1; Pred. No.
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Pred. No. 29
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95;
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1493 CAGACTTC 1500

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US-09-990-186-456/c
Sequence 456, Application US/09990186
Publication No. US20030068675A1
GENERAL INFORMATION:
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US-09-990-186-455/c
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US-09-989-789-456/c
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                                                                           RESULT 33
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TITLE OF INVENTION: POSITION DEPENDENT RECOGNITION
TITLE OF INVENTION: TRIPLETS BY ZINC FINGERS
FILE REFERENCE: 8325-0011.20 / S11-US2
CURRENT APPLICATION NUMBER: US/09/989,789
CURRENT FILING DATE: 2002-03-25
NUMBER OF SEQ ID NOS: 4085
SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 456
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Matches
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Best Local Similarity
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APPLICANT: LIU, Qiang
                                                                                                                                                                                                                                                                                                                                LENGTH: 9
TYPE: DNA
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Mismatches
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Sequence 456, Application US/09989994
Publication No. US20030104526A1
GENERAL INFORMATION:
APPLICANT: LIU, Qiang
TITLE OF INVENTION: POSITION DEPENDENT RECOGNITION OF
TITLE OF INVENTION: TRIPLETS BY ZINC FINGERS
FILE REFERENCE: 8325-0011.20 / S11.US2
CURRENT APPLICATION NUMBER: US/09/989,994
CURRENT FILING DATE: 2001-11-20
NUMBER OF SEQ ID NOS: 4085
SOFTWARE: Patentin Ver: 2.0
SEQ ID NO 456
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US-09-990-186-456
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US-09-989-994-456/c
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Best Local S
Matches 8
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TITLE OF INVENTION: TRIPLETS BY ZINC FINGERS
FILE REFERENCE: 8325-0011.21 / S11-US3
CURRENT APPLICATION NUMBER: US/09/990,186
CURRENT FILING DATE: 2001-11-20
NUMBER OF SEQ ID NOS: 4085
SOFTWARE: PATENTIN Ver. 2.0
SEQ ID NO 456
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LENGTH: 9
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Best Local
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TITLE OF INVENTION: POSITION DEPENDENT RECOGNITION OF
TITLE OF INVENTION: TRIPLETS BY ZINC FINGERS
FILE REFERENCE: 8325-0011.20 / S11-US2
CURRENT APPLICATION NUMBER: US/09/989,994
CURRENT FILING DATE: 2001-11-20
NUMBER OF SEQ ID NOS: 4085
SOFTWARE: Patentin Ver. 2.0
                  TYPE: D
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ORGANISM: Artificial Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                   26.7%;
                                                                                                                                                                                                                                                                                                                                                                                                                      0
                                                                                                                                                                                                                                                                                                                                                                                                                                     Score 8;
Pred. No.
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Query Match
Best Local Similarity
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               SEQ ID NO 6
LENGTH: 9
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Matches 8; Conserv
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                                                                         TITLE OF INVENTION: METHOD AND KIT FOR THE DETERMINATION OF TITLE OF INVENTION: CELLULAR ACTIVATION PROFILES FILE REFERENCE: VANM212.001CP1
CURRENT APPLICATION NUMBER: US/10/339,161
CURRENT FILING DATE: 2003-01-07
PRIOR APPLICATION NUMBER: US 09/816,763
PRIOR APPLICATION NUMBER: US 09/816,763
PRIOR FILING DATE: 2001-03-23
PRIOR FILING DATE: 2001-03-24
NUMBER OF SEQ ID NOS: 27
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: Fang, Yu
APPLICANT: Wang, Xiao-Yang
APPLICANT: Wang, Xiao-Yang
APPLICANT: Turpin, Pierre
TITLE OF INVENTION: Methods of detecting multiple DNA
TITLE OF INVENTION: binding protein and DNA interactions in a sample,
TITLE OF INVENTION: devices, systems and kits for practicing the same.
FILE REFERENCE: CLON-071
CURRENT APPLICATION NUMBER: US/10/113,877
CURRENT FILING DATE: 2002-03-29
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     PRIOR APPLICATION NUMBER: 60/314,330
PRIOR FILING DATE: 2001-08-20
NUMBER OF SEQ ID NOS: 192
SOFTWARE: FastSEQ for Windows Version 4.0
                                                            SOFTWARE: FastSEQ for Windows Version 4.0
                                                                                                                                                                                                                                                                                             APPLICANT: Remacle, Jose
APPLICANT: Renard, Patricia
APPLICANT: Art, Muriel
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          PRIOR APPLICATION NUMBER: 60/280,658 PRIOR FILING DATE: 2001-03-30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TYPE: DNA
ORGANISM: Artificial Sequence
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TYPE: DNA
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100.0%; Pred. No. 95;
tive 0; Mismatches
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; FEATURE: ; OTHER INFORMATION: Transcription factor SMAD US-10-339-161-6
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US-10-277-494-147
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US-10-277-494-147/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      RESULT 39
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: Ribozyme Pharmaceuticals, Inc.
APPLICANT: McSwiggen, Jim
TITLE OF INVENTION: Nucleic Acid Treatment of Diseases or conditions Related
TITLE OF INVENTION: Epidermal Growth Factor Receptors
FILE REFERENCE: MSHB00-958-K (400/064)
CURRENT APPLICATION NUMBER: US/10/277,494
CURRENT FILING DATE: 2002-10-21
RUMBER OF SEQ ID NOS: 446
SOFTWARE: PatentIn version 3.0
SEQ ID NO 147
TYPE PART
                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 9, Application US/08463404 Publication No. US20020127634A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 147, Application US/10277494 Publication No. US20033186909A1 GENERAL INFORMATION:
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Matches 8; Conserv
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CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/463,404
FILING DATE: 05-JUN-1995
                                                                                                     ZIP: 90071-2066
COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: storage
                                                                                                                                                                                                                                                             CORRESPONDENCE ADDRESS:
ADDRESSEE: Lyon & Ly
                                                                                                                                                                                                                                                                                                               TITLE OF INVENTION: TITLE OF INVENTION:
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                                                                                                                                                                                                                                                                                               NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                  APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT:
                                                                                                                                                                                     ADDRESSEE: Lyon & Lyon
STREET: 633 West Fifth Street
STREET: Suite 4700
CITY: Los Angeles
STATE: California
                                                   COMPUTER: IBM Compatible OPERATING SYSTEM: IBM P.C. SOFTWARE: Word Perfect 5.1
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                                                                                                                                                                       COUNTRY:
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milarity 100.0%;
Conservative 0
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Woodring E. Wright
Elizabeth Blackburn
                                                                                                                                                                           U.S.A.
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Pred. No
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Pred. No. 95;
                                                                       DOS 5.0
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Levels

PRIOR APPLICATION DATA:

08/060,952

CLASSIFICATION: 435 APPLICATION NUMBER:

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GENERAL INFORMATION:
APPLICANT: GENZYME CORPORATION
APPLICANT: ROBERTS, BRUCE
APPLICANT: SHANKARA, SRINIVAS
TITLE OF INVENTION: PREPARATION AND USE OF SUPERIOR VACCINES
FILE REFERENCE: GAO201C
CURRENT APPLICATION NUMBER: US/10/033,145
CURRENT FILING DATE: 2001-11-05
PRIOR APPLICATION NUMBER: FCT/US99/13800
PRIOR APPLICATION NUMBER: FCT/US99/13800
PRIOR FILING DATE: 1999-06-18
NUMBER OF SEQ ID NOS: 2137
SOFTMARE: PATENTIN VERSION 3.0
SEQ ID NO 185
LENGTH: 10
TYPE: DNA
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US-10-033-145-185/c
                                                                                                                                RESULT 41
US-10-033-145-1011
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                                                                         Sequence 1011, Application US/10033145
Publication No. US20020151515A1
GENERAL INFORMATION:
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Best Local :
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                     APPLICANT: GENZYME CORPORATION APPLICANT: ROBERTS, BRUCE APPLICANT: SHANKARA, SRINIVAS
APPLICANT: SHANKARA, SRINIVAS
TITLE OF INVENTION: PREPARATION AND USE OF SUPERIOR VACCINES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           FILING DATE: May 13, 1993
APPLICATION NUMBER: 07/882,438
FILING DATE: May 13, 1992
APPLICATION NUMBER: 08/038,766
FILING DATE: March 24, 1993
ATTORNEY/ACENT IMPORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SEQUENCE CHARACTERISTICS:
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REGISTRATION NUMBER: 32,327
REFERENCE/DOCKET NUMBER: 20:
TELECOMMUNICATION INFORMATION:
TELEPHONE: (213) 489-1600
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Local Similarity 100.0%; I
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0; Mismatches
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Mismatches
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; TYPE: DNA
; ORGANISM: Homo sapiens
US-10-033-145-1011
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              US-10-113-030-3/c
                                                                                                                                                                                                                                                                                                                             US-10-358-818-3/c
Sequence 3, Application US/10358818
Publication No. US20030162213A1
GENERAL INFORMATION:
APPLICANT: Fuller, Carl
APPLICANT: Sood, Anup
APPLICANT: Sood, Anup
APPLICANT: Nelson, John
TITLE OF INVENTION: Terminal-Phosphate-Labeled Nucleotides and Methods of Use
FILE REFERENCE: PB0156-1CIP
CURRENT APPLICATION NUMBER: US/10/358,818
CURRENT FILING DATE: 2003-02-05
PRIOR APPLICATION NUMBER: US 60/315,798
PRIOR APPLICATION NUMBER: US 60/315,798
PRIOR APPLICATION NUMBER: US 10/113,030
PRIOR FILING DATE: 2002-04-01
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: Fuller, Carl
APPLICANT: Sood, Anup
APPLICANT: Sood, Anup
APPLICANT: Kumar, Shiv
TITLE OF INVENTION: Terminal-Phosphate-Labeled Nucleotides and Methods of Use
FILE REFERENCE: PB0156-1
CURRENT APPLICATION NUMBER: US/10/113,030
CURRENT FILING DATE: 2002-04-01
PRIOR APPLICATION NUMBER: US 60/315,798
PRIOR APPLICATION NUMBER: US 60/315,798
PRIOR APPLICATION NUMBER: US 60/315,798
PRIOR PRIOR FILING DATE: 2001-08-29
NUMBER OF SEQ ID NOS: 3
SOFTWARE: Patentin version 3.1
SEQ ID NO SEQ ID NOS: 3
SOFTWARE: Patentin version 3.1
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Publication No. US20030077610A1
GENERAL INFORMATION:
APPLICANT: Nelson, John
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Best Local Similarity luv.
Thickes 8; Conservative
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Matches 8; Conserv
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PRIOR FILING DATE: 1999-06-18
NUMBER OF SEQ ID NOS: 2137
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CURRENT APPLICATION NUMBER: US/10/033,145
CURRENT FILING DATE: 2001-11-05
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TYPE: DNA ORGANISM: Artificial Sequence
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100.0%; Pred. No
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Sequence 344, Application US/10330627

Publication No. US20030175771A1

GENERAL INFORMATION:

APPLICANT: Velculescu, Victor E.

APPLICANT: Velculescu, Victor E.

APPLICANT: Vegelstein, Bert

TITLE OF INVENTION: Human Transcriptomes

FILE REFERENCE: 001107.00319

CURRENT APPLICATION NUMBER: US/10/330,627

CURRENT APPLICATION NUMBER: US 09/448,480

PRIOR APPLICATION NUMBER: US 09/448,480

PRIOR APPLICATION NUMBER: US 09/448,480

PRIOR FILING DATE: 1999-11-24

NUMBER OF SEQ ID NOS: 1564

SOFTMARE: FastSEQ for Windows Version 4.0

SEQ ID NO 344

LENGTH: 10

TYPE: DNA

ORGANISM: Homo Sapiens

US-10-330-627-344
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ; NUMBER OF SEQ ID NOS: 3
; SOFTWARE: PatentIn version 3.1
; SEQ ID NO 3
; LENGTH: 10
TYPE: DNA
; ORGANISM: Artificial Sequence
; FEATURE:
; OTHER INFORMATION: DNA Template
US-10-358-818-3
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US-10-330-627-344/c
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Search completed: April 15, 2004, 16:38:43 Job time : 0.001 secs
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RESULT 1
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Listing first 1 summaries
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HOA28-1-G6 HOA (Human Osteoarthritic Cartilage) Homo sapiens cDNA, mRNA sequence.
GlaxoSmithKline
709 Swedeland Road, P.O. Box 1539, King of Prussia,
Tel: 610-270-7245
                                                                                                                                   Identification and initial characterization of 5000 expressed sequenced tags (ESTs) each from adult human normal and osteoarthritic cartilage cDNA libraries Osteoarthr. Cartil. 9 (7), 641-653 (2001)
                                                                                                                                                                                                                  Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.

1 (bases 1 to 11)
Kumar, S., Connor, J.R., Dodds, R.A., Halsey, W., Van Horn, M., Mao, J., Sathe, G., Mui, P., Agarwal, P., Badger, A.M., Lee, J.C., Gowen, M. and Lark, M.W.
                                                           UW2109
                                                                            Contact: Sanjay Kumar
                                                                                                                     21482651
                                                                                                                                                                                                                                                                                                                                       Homo sapiens
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Email: sanjay_kumar-l@gsk.com
Seq primer: T7.
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                                                                                                                 Conservative
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/db_xref="taxon:9606"
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